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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
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A Geneseq_16Dec04:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2003s:*
6: geneseqp2003bs:*
7: geneseqp2003bs:*
8: geneseqp2004s:*
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Gapop 10.0 , Gapext 0.5
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329
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 YPVRCLLPPATGPCKARIIR......RGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                     2105692 seqs, 386760381 residues
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match L	Length	DB	ID	Description
_ ;	329	100.0	58	٥	ABG71914	Abg71914 Human Kun
N	326	99.1	82	δ	ABG71915	71915 Human
u	314	95.4	58	σ	ABG71917	Human
4.	314	95.4	111	σ	ABG71920	Human
5	291	88.4	58	σ	ABG71916	Human
,6	291	8.4		σ	ABG71919	Human
<u>\</u>	264	80.2		4	AAB60623	Human
8	264			σ	ABG71912	2 Human
9	264	80.2	58	8	ADL16831	.6831 Human
10	264	80.2	111	δ	ABG71918	Human
11	264	80.2	189	44	AAB60634	Aab60634 Human pro
12	264	80.2	189	œ	ADL16845	Adl16845 BTL.010 p
13	. 264		1235	æ	ADO55146	Ado55146 Protein #
14	264	80.2	1280	σ	ABR58408	Abr58408 Human NOV
15	220.5	67.0	51	4.	AAB60631	
16	220.5	67.0	51	œ	ADL16839	Adl16839 Kunitz ty
17	208	63.2	109	N	AAR22985	Aar22985 Aprotinin
18	203	61.7	111	N	AAR22979	
19	202	61.4	109	N	AAR22983	Aar22983 Aprotinin
20	202	61.4	128	N	AAW82770	Aaw82770 Bovine bi
21	199	60.5	57	N	AAW47433	Aaw47433 Aprotinin
22	198		141	N	AAR22982	Aar22982 Aprotinin
23	195	59.3	128	N	AAW82769	
24	194	59.0	111	N	AAR22980	
25	194	59.0	128	N	AAW82768	Aaw82768 Bovine bi

F 31-MAY-2002; 2002WO-DK000372. X 31-MAY-2001; 2001DK-00000859. R 05-JUL-2001; 2001US-0303180P.	D 05-DEC-2002. X	odifference	c-difference 1	c-difference 9 c-difference 1	sapiens.	KW Human; protease inhibitor; kunitz; HKI-18 KW anticoagulant; coagulant; cardiant; CBP;] KW systemic inflammatory response syndrome; KW shock syndrome; disseminated intravascula; KW hyperfibrinolytic haemorrhage; myocardial KW hyperfibrinolytic haemorrhage; myocardial KW cardiopulmonary bypass-induced pulmonary allergy-induced protease release; deep ve. KW adult respiratory distress syndrome; chrol XX	an Kunitz protease inhibitor	22-JAN-2003 (first entry)	71914	SULT 1 371914 ABG71914 standard; protein; 58 AA.	ALIGNMENTS	41 192 58.4 58 2 AAR891477 42 192 58.4 58 2 AAR64119 43 192 58.4 58 2 AAR64119 44 192 58.4 58 2 AAR66410 44 192 58.4 58 2 AAR7691983 45 192 58.4 58 2 AAR79983	192 58.4 58 2 192 58.4 58 2	192 58.4 58 1 192 58.4 58 2	193 58.7 141 2 193 58.7 141 2	193 58.7 128 2 193 58.7 128 2	193 58.7 57 2	193 58.7 57 2 193 58.7 57 2	193 58.7 57 2	194 59.0 141 2	194 59.0 141 2
		p substituted by Val" s substituted by Arg"	r substituted by Pro" a-Asp-Trp-Ala-Ala substituted by Lys-	r substituted by Pro" s substituted by Thr"		unitz; HKI-18; antiinflammatory; rdiant; CBP; psoriasis; emphysema; se syndrome; acute pancreatitis; intravascular coagulation; mutein; e; myocardial infarction; mutein; ed pulmonary injury; rheumatoid arthritis; ease; deep vein thrombosis; yndrome; chronic inflammatory bowel disease.	protein HKI-18, mutant #1.				MENTS	Adroido macure Aarolia Aprocit Aarolia Human I Aaw64110 Cationi Aaw79983 Bovine	Aar47179 Aar78426	Aap93401 Aar27373		Aaw82771 Aaw82772	Aaw47435	Aaw47434 Aaw47432	Aaw47436	Aar22986	Aar22988 Aprotini

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         /note= "Wild-type Ala-Asp-Trp-Ala-Ala substituted by Arg-Ala-Arg-Ile-Ile"
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Best Local Similarity
Matches 57; Conserv
Human; protease inhibitor; kunitz; HKI-18; antiinflammatory; anticoagulant; coagulant; cardiant; CBP; psoriasis; emphysema; systemic inflammatory response syndrome; acute pancreatitis; shock syndrome; disseminated intravascular coagulation; mutant; hyperfibrinolytic haemorrhage; myocardial infarction; mutein; cardiopulmonary bypass-induced pulmonary injury; rheumatoid arthritis; allergy-induced protease release; deep vein thrombosis; adult respiratory distress syndrome; chronic inflammatory bowel disease
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Misc-difference 39
                                                                                                                                                             Human Kunitz protease inhibitor protein HKI-18, mutant
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                                                                                                                                                                                                                                                                                                                                                                                                                                          acid sequence being a generic sequence for a Kunitz-type protease inhibitor or a variant where the sequence is at least 80% identical to the kunitz domain of human kunitz type protease inhibitor HKI-18. Also included are an isolated polypeptide obtainable by cultivation of a host cell comprising a polynucleotide construct encoding kunitz-type protein in an appropriate growth medium under conditions allowing expression of
               ABG71920;
                                           ABG71920 standard;
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jorgensen MU,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAY-2002; 2002WO-DK000372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                inflammatory
human HKI-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (NOVO ) NOVO NORDISK AS
                                                                                                                                                                                          Local
                                                                                                                                                                             55;
                                                                                                                                                                                            Similarity
                                                                                                                                   YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                      58
                                                                                                                YPVRCLLPPATGPCKARAARWYFVASVGQCNRFVYGGCHGNANNFASEQECMSSCQGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protease inhibitor used
                                                                                                                                                                         95.4%;
llarity 94.8%;
Conservative
                                                                                                                                                                                                                                      Š
                                                                                                                                                                                                                                                                 mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001DK-00000859
2001US-0303180P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note=
34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note=
15. .17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "Wild-type Trp substituted by Val"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note= "Wild-type His substituted by Thr"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note= "Wild-type Ser substituted"
                                         protein; 111
                                                                                                                                                                                                                                                                sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ά
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  . 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Wild-type Ser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Wild-type Ala-Asp-Trp substituted by Lys-Ala-Arg"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Olsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English
                                                                                                                                                                           0,
                                                                                                                                                                        Score 314; DB 6;
Pred. No. 6.6e-30;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ,
H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Petersen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            substituted by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the treatment of inflammatory
                                                                                                                                                                                                     Length 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ş
                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pro"
                                                                                                                                                                           0
                                                                                                                  58
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The invention relates to an isolated polypeptide comprising a 51 amino CC acid sequence being a generic sequence for a Kunitz-type protease CC inhibitor or a variant where the sequence is at least 80% identical to CC the kunitz domain of human Kunitz type protease inhibitor HKI-B. Also CC included are an isolated polypeptide obtainable by cultivation of a host CC cell comprising a polynucleotide construct encoding kunitz-type protein in an appropriate growth medium under conditions allowing expression of CC the polynucleotide construct encoding the kunitz-type protein for the polynucleotide construct encoding the kunitz-type CC culture medium, a polynucleotide construct encoding the kunitz-type CC protein and a host cell comprising the polynucleotide construct. The CC kunitz-type protein is used for the preparation of a medicament for the CC kunitz-type protein is used for the preparation, acute pancreatitis, CC shock syndrome, disseminated intravascular coagulation, hyperfibrinolytic haemorrhage, myocardial infarction, for prevention of blood loss during CC major surgery, cardiopulmonary bypass (CPB)-induced pulmonary injury, CC allergy-induced protease release, deep vein thrombosis, emphysema, Crimanatory bowel disease, and psoriasis. The present sequence CC inflammatory bowel disease, and psoriasis. The present sequence CC represents a mutant human HKI-18 with a yeast 22L signal peptide and a CC KEX-2 cleavage site, expressed in yeast cells
                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Fig 8; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kunitz-type protease inhibitor used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jorgensen MU,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-2001; 2001DK-00000859.
05-JUL-2001; 2001US-0303180P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-MAY-2002; 2002WO-DK000372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cardiopulmonary bypass-induced pulmonary injury; rheumatoid arthritis; allergy-induced protease release; deep vein thrombosis; adult respiratory distress syndrome; chronic inflammatory bowel disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       anticoagulant; coagulant; CBP; psoriasis; emphysema; mutein; systemic inflammatory response syndrome; acute pancreatitis; shock syndrome; disseminated intravascular coagulation; yeast; hyperfibrinolytic haemorrhage; myocardial infarction; 2121;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-041403/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200296938-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cleavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NOVO ) NOVO NORDISK AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            212L-HKI-18-2 fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protease inhibitor; kunitz; HKI-18; antiinflammatory; mutant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                68.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= 212L_signal_peptide
52. .53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "Wild-type Ala-Asp-Trp substituted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label= KEX2_cleavage_site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "Wild-type His substituted by Thr"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in the treatment of inflammatory
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RESULT 5
ABG71916
ID ABG7
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Matches
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The invention relates to an isolated polypeptide comprising a 51 amino acid sequence being a generic sequence for a Kunitz-type protease inhibitor or a variant where the sequence is at least 80% identical to the kunitz domain of human kunitz type protease inhibitor HKI-18. Also included are an isolated polypeptide obtainable by cultivation of a host cell comprising a polymucleotide construct encoding kunitz-type protein in an appropriate growth medium under conditions allowing expression of the polynucleotide construct and recovering the polypeptide from the culture medium, a polymucleotide construct encoding the kunitz-type protein and a host cell comprising the polynucleotide construct. The kunitz-type protein is used for the preparation of a medicament for the treatment of systemic inflammatory response syndrome, acute pancreatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAY-2001; 2001DK-00000859
05-JUL-2001; 2001US-0303180P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                      Claim 20; Page 49; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kunitz-type protease inhibitor used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-041403/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-MAY-2002; 2002WO-DK000372
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jorgensen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human Kunitz protease inhibitor protein HKI-18, mutant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; protease inhibitor; kunitz;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          respiratory distress syndrome; chronic inflammatory bowel disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Olsen OH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   58
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Pred. No. 1.3e-29;
0; Mismatches 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the treatment of inflammatory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        γģ
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RESULT 6
ABG71319
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XX Huma
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Best Local S
Matches 52
                                                     WPI; 2003-041403/03
                                                                                                                                                                                                               31-MAY-2001; 2001DK-00000859.
05-JUL-2001; 2001US-0303180P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   shock syndrome, disseminated intravascular coagulation, hyperfibrinolytic haemorrhage, myocardial infarction, for prevention of blood loss during major surgery, cardiopulmonary byges (CPB)-induced pulmonary injury, allergy-induced protease release, deep vein thrombosis, emphysema, rheumatoid arthritis, adult respiratory distress syndrome, chronic inflammatory bowel disease, and psoriasis. The present sequence is a
                                                                                                                                                                                                                                                                                             31-MAY-2002; 2002WO-DK000372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; protease inhibitor; kunitz; HKI-18; antiinflammatory; mutant; anticoagulant; coagulant; cardiant; CDP; psoriasis; emphysema; mutein; systemic inflammatory response syndrome; acute pancreatitis; shock syndrome; disseminated intravascular coagulation; yeast; shock syndrome; disseminated intravascular coagulation; 212L; hyperfibrinolytic haemorrhage; myocardial infarction; 212L; cardiopulmonary bypass-induced pulmonary injury; rheumatoid arthritis; allergy-induced protease release; deep vein thrombosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABG71919 standard;
                                                                                                       Jorgensen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cleavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human 212L-HKI-18-1 fusion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABG71919;
                                                                                                                                                            (NOVO ) NOVO
                                                                                                                                                                                                                                                                                                                                                                                                        WO200296938-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            respiratory distress
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     YPVRCLLPPATGSCKAWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS
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                                                                                                                                                            NORDISK AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                    Bang S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /label= KEX2_cleavage_site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "Wild-type Ser substituted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /label= 212L_signal_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               note= "Wild-type His substituted by Thr"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 label= HKI_18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      . 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Wild-type Ala-Asp substituted by Lys-Ala"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            syndrome; chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 291; DB 6;
Pred. No. 3.7e-27;
                                                                                                       읝
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                                                                                                       Petersen LC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammatory bowel disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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Kunitz-type protease inhibitor used

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treatment

of inflammatory

disorders Example

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8; 52pp; English.

of.

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RESULT 7

AAB60623

ID AAB6

XX AAB6

XX AAB6

XX AAB6

XX AAB6

XX Huma

XX Huma

XX Huma

XX Huma

XX Homo

XX Pat

XX OS-J

XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              shock syndrome, disseminated intravascular coagulation, hyperfibrinolytic haemorrhage, myocardial infarction, for prevention of blood loss during major surgery, cardiopulmonary bypass (CPB) induced pulmonary injury, allergy-induced protease release, deep vein thrombosis, emphysema, rheumatoid arthritis, adult respiratory distress syndrome, chronic inflammatory bowel disease, and psoriasis. The present sequence represents a mutant human HKI-18 with a yeast 2121 signal peptide and a KEX-2 cleavage site, expressed in yeast cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            acid sequence being a generic sequence for a Kunitz-type protease inhibitor or a variant where the sequence is at least 80% identical to the kunitz domain of human kunitz type protease inhibitor HKI-18. Also included are an isolated polypeptide obtainable by cultivation of a host cell comprising a polynucleotide construct encoding kunitz-type protein in an appropriate growth medium under conditions allowing expression of the polynucleotide construct and recovering the polypeptide from the culture medium, a polynucleotide construct encoding the kunitz-type protein and a host cell comprising the polynucleotide construct. The kunitz-type protein is used for the preparation of a medicament for the kunitz-type protein is used for the preparation of a medicament for the kunitz-type systemic inflammatory response syndrome, acute pancreatitis, shock syndrome, acute pancreatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           proteinase-3 inhibitor; Kunitz domain; emphysema; idiopathic pulmonary fibrosis; adult respiratory distress syndrome; cystic fibrosis; rheumatoid arthritis; organ failure; glomerulonephritis; platelet activation; blood coagulation; neutrophil activation; monocyte activation; angioplasty; inflammatory disease; lung injury;
                                  N-PSDB; AAF59750
                                                                  WPI; 2001-190860/19
                                                                                                                                Davies C,
                                                                                                                                                                                               (DAVI/) DAVIES C.
(CHEN/) CHEN D.
(ROCZ/) ROCZNIAK S.
                                                                                                                                                                                                                                                                                                                                                                                                     05-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human BTL.010; neutral serine protease inhibitor; elastase inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-APR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB60623;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB60623 standard; protein; 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 111
                                                                                                                                                                                                                                                                                                                                      05-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US6180607-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vascular injury; nephrotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated polypeptide comprising a 51 amino acid sequence being a generic sequence for a Kunitz-type protease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  54 YPVRCLLPPATGSCKAWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS
                                                                                                                                Chen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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A
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                                                                                                                                                                                                                                                                                                                                                                                                         99US-00369494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inhibitor BTL.010 Kunitz domain,
                                                                                                                                    Roczniak S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  88.4%;
89.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 291; DB 6;
Pred. No. 7.2e-27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antirheumatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antiarthritic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO:1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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WO200296938-A2

5. .55
/label= Kunitz_domain Location/Qualifiers

/note= "This domain is

claimed in claim

Homo sapiens.

cardiopulmonary bypass-induced pulmonary injury; rheumatoid arthritis; allergy-induced protease release; deep vein thrombosis;

chronic

inflammatory bowel disease

respiratory distress syndrome;

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ABG7191
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AC ABB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; protease inhibitor; kunitz; HKI-18; antiinflammatory; anticoagulant; coagulant; cardiant; CBP; psoriasis; emphysema; systemic inflammatory response syndrome; acute pancreatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel human serine protease inhibitor of the Kunitz family, BTL.010 (fragments given in AAB60623, AAB60631 and AAB60634). The BTL.010 protein is thought to preferentially inhibit neutral serine proteases such as elastase and proteinase-3, relative to trypsin-like and chymotrypisn-like proteases A substantial proportion of the BTL.010 protein Kunitz domain (AAB60631) was identified via homology searching in the GenBank high throughput genomic (HTG) DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                            shock syndrome; disseminated intravascular coagulation;
hyperfibrinolytic haemorrhage; myocardial infarction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Kunitz protease inhibitor protein HKI-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABG71912 standard; protein; 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel serine proteinase inhibitor of the Kunitz family, BTL.010 useful for treating emphysema, cystic fibrosis, adult respiratory distress syndrome, rheumatoid arthritis, organ failure and glomerulonephritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ьосат
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48; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    80.2%;
ilarity 82.8%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 264; DB 4;
Pred. No. 6.3e-24;
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RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to an isolated polypeptide comprising a 51 amino CC acid sequence being a generic sequence for a Kunitz-type procease CC inhibitor or a variant where the sequence is at least 80% identical to the Kunitz domain of human Kunitz type protease inhibitor HKI-18. Also CC included are an isolated polypeptide obtainable by cultivation of a host cell comprising a polynucleotide construct encoding kunitz-type protean cC in an appropriate growth medium under conditions allowing expression of the polynucleotide construct encoding the kunitz-type protein for the polynucleotide construct encoding the kunitz-type culture medium, a polynucleotide construct encoding the kunitz-type protein and a host cell comprising the polynucleotide construct. The cunitz-type protein is used for the preparation of a medicament for the creatment of systemic inflammatory response syndrome, acute pancreatitis, shock syndrome, disseminated intravascular coagulation, hyperfibrinolytic haemorrhage, myocardial infarction, for prevention of blood loss during cmajor surgery, cardiopulmonary bypass (CPB)-induced pulmonary injury, callergy-induced protease release, deep vein thrombosis, emphysema, creating and partities, adult respiratory distress syndrome, chronic inflammatory bowel disease, and psoriasis. The present sequence is human continuation action and proteins and present sequence is human continuation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                             BTL.010 protein; serine proteinase inhibitor; Kunitz domain; emphysema; idiopathic pulmonary fibrosis; adult respiratory distress syndrome; cystic fibrosis; rheumatoid arthritis; organ failure; glomerulonephritis; inflammatory disease; oxidative burst; platelet activation;
                                    10-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                     Human BTL.010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADL16831;
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                                                                                               US6689582-B1.
                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADL16831 standard;
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05-JUL-2001; 2001US-0303180P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NOVO ) NOVO
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                                                                                                                                                                                                                  coagulation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                              protein Kunitz domain (KD) peptide.
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                                                                                                                                                                                                                         therapy; human;
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Pred. No. 6.3e-24;
0; Mismatches 10
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Best Local S
Matches 48
                                                                                                                                                                                                                                                                                                                                                                                                      hyperfibrinolytic haemorrhage; myocardial infarction; 212L; cardiopulmonary bypass induced pulmonary injury; rheumatoid arthritis; allergy-induced protease release; deep vein thrombosis; adult respiratory distress syndrome; chronic inflammatory bowel disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the Kunitz family and its corresponding nucleic acid sequence. The sequences of the invention are useful for treating diseases, e.g. emphysema, idiopathic pulmonary fibrosis, adult reagiratory distress syndrome, cystic fibrosis, rheumatoid arthritis, organ failure, glomerulonephritis or inflammatory diseases. The BTL.010 protein is also useful for preventing neutrophil and monocyte activation and formation of active oxygen species during the oxidative burst of stimulated granulocytes. It is also useful for reducing platelet activation and blood coagulation. BTL.010 DNA is useful in gene thorapy. The present sequence is human BTL.010 protein Kunitz domain (KD) peptide.
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                                                                                                                                                                                                                                                                                                                                              Saccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           anticoagulant; coagulant; cardiant; CBP; psoriasis; emphysema; systemic inflammatory response syndrome; acute pancreatitis; shock syndrome; disseminated intravascular coagulation; yeast;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; protease inhibitor; kunitz; HKI-18; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human HKI-18/212L signal peptide.
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                                                                                                                                                                                                                                                                                        Chimeric.
                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Davies C,
                                                                                                                                                                                                Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to BTL.010, a human serine proteinase inhibitor of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1;
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                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                       /label= 212L_signal_peptide
                                                     /label= HKI_18
                                                                                                       /label= KEX2_cleavage_site
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Pred. No. 6.3e-24;
0; Mismatches 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                   disease
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WO200296938-A2

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VRESULT 11
AAB60634
ID AAB60
XX Human
XX Homo
XX Homo
XX Homo
XX Homo
XX Homo
XX US618
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC inhibitor or a variant where the sequence is at least 80% identical to CC the kunitz domain of human kunitz type protease inhibitor HKI-18. Also CC included are an isolated polypeptide obtainable by cultivation of a host Cell comprising a polynuclocide construct encoding kunitz-type protein CC in an appropriate growth medium under conditions allowing expression of CC the polynucleotide construct and recovering the polypeptide from the CC culture medium, a polynucleotide construct encoding the kunitz-type cC protein and a host cell comprising the polynucleotide construct. The CC kunitz-type protein is used for the preparation of a medicament for the creatment of systemic inflammatory response syndrome, acute pancreatitis, CC shock syndrome, disseminated intravascular coagulation, hyperfibrinolytic haemorrhage, myocardial infarction, for prevention of blood loss during CC major surgery, cardiopulmonary bypass (CPB) induced pulmonary injury, CC allergy-induced protease release, deep vein thrombosis, emphysema, CC inflammatory bowel disease, and psoriasis. The present sequence CC represents human HKI-18 with a yeast 121 signal peptide and a KEX-2 class of the present sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 48
                                                                        Human BTL.010; neutral serine protease inhibitor; elastase inhibitor; proteinase-3 inhibitor; Kunitz domain; emphysema; idiopathic pulmonary fibrosis; adult respiratory distress syndrome; cystic fibrosis; rheumatoid arthritis; organ failure; glomerulonephritis; platelet activation; blood coagulation; meutrophil activation; angioplasty; inflammatory disease; lung injury; vascular injury; nephrotropic; antirheumatic; antiarthritic.
                                                                                                                                                                                                                                    Human protease inhibitor BTL.010 ORF protein sequence, SEQ ID NO:15.
                                                                                                                                                                                                                                                                                                                     AAB60634;
                                                                                                                                                                                                                                                                                                                                                             AAB60634 standard; protein; 189 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated polypeptide comprising a 51 acid sequence being a generic sequence for a Kunitz-type protease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Fig 7; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jorgensen MU,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-MAY-2001; 2001DK-00000859
05-JUL-2001; 2001US-0303180P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-MAY-2002; 2002WO-DK000372.
                                     Homo sapiens
                                                                                                                                                                                                                                                                              27-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cleavage site, expressed in yeast cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kunitz-type protease inhibitor used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2003-041403/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                               54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                               YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B
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                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          80.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Olsen OH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 264; DB 6;
Pred. No. 1.2e-23;
0; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Petersen LC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in the treatment of inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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RESULT 12
ADL16945
ID ADL16
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AC ADL16
XZ
AC ADL16
XZ
DT 06-MA
XX
DE BTL.0
XX
KW BTL.0
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                                                                                                                                                                                                                                                                                                                                                                                                                              neutral serine proteases such as elastase and proteinase-3, relative to trypsin-like and chymotrypisn-like proteases. A substantial proportion of the BTL.010 protein Kunitz domain (AAB60631) was identified via homology searching in the GenBank high throughput genomic (HTG) DNA sequence database using the Kunitz domain sequences AAB60630, and was confirmed as being novel using the Kunitz domain sequences AAB60630, and was confirmed as Deing novel using the Kunitz domain sequences AAB60632, and AAB60633.

This sequence information was extended to provide a larger region of BTL.010 protein sequence data (AAB60634) by identifying an open reading frame (ORF) which comprised DNA encoding the BTL.010 Kunitz domain corresponding to bases 16016-16414 of GenBank accession number AC004846.

The entire BTL.010 Kunitz domain sequence (AAB60623) was obtained from the BTL.010 ORF-encoded sequence. The BTL.010 protein, and pharmaceutical compositions comprising it, may be used for inhibiting protease activity particularly that of leukocyte elastase, in the prevention, treatment or ameliotation of medical conditions such as emphysema, idiopathic pulmonary fibrosis, adult respiratory distress syndrome, cystic fibrosis, rheumatoid arthritis, organ failure or glomerulomephritis BTL.010 condition such as platelet activation, blood coagulation, neutrophil activation, or monocyte activation. BTL.010 is also useful for the
                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                      BTL.010 protein related protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel serine proteinase inhibitor of the Kunitz family, BTL.010 useful for treating emphysema, cystic fibrosis, adult respiratory distress syndrome, rheumatoid arthritis, organ failure and glomerulonephritis.
                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 189 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prophylactic or therapeutic treatment of patients undergoing angioplasty, and for the treatment of inflammatory diseases and diseases involving lung and vascular injury. The present sequence represents a portion of the human BTL.010 protease inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel human serine protease inhibitor of Kunitz family, BTL.010 (fragments given in AAB60623, AAB60631 and AAB60634). The BTL.010 protein is thought to preferentially inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Col 7-8; 17pp;
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                                                                                                                                                            ADL16845 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Davies C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (DAVI/) DAVIES C.
(CHEN/) CHEN D.
(ROCZ/) ROCZNIAK S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001-190860/19
                                                                                                                                                                                                                                                                                                     1 YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                              48;
                                                                                                                                                                                                                                                               YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chen D,
                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-00369494
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                                                                                                                                                            protein;
                                                                                                                                                                                                                                                                                                                                                                 80.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Roczniak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                              <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                 Score 264; DB 4;
Pred. No. 2.1e-23;
                                        #2
                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                              10;
                                                                                                                                                                                                                                                                                                                                                                                      Length 189;
                                                                                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                              Gaps
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BTL.010 protein; serine proteinase inhibitor; Kunitz family; emphysema;

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ADO55146
ID ADO55
XX ADO55
XX ADO55
XX ADO55
XX ADO55
XX ADO55
XX CDT 15-JU
DT 15-JU
DT 15-JU
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Best Local Similarity
Matches 48; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to BTL.010, a human serine proteinase inhibitor of the Kunitz family and its corresponding nucleic acid sequence. The sequences of the invention are useful for treating diseases, e.g. emphysema, idiopathic pulmonary fibrosis, adult respiratory distress syndrome, cystic fibrosis, rheumatoid arthritis, organ failure, glomerulonephritis or inflammatory diseases. The BTL.010 protein is also useful for preventing neutrophil and monocyte activation and formation of active oxygen species during the oxidative burst of stimulated granulocytes. It is also useful for reducing platelet activation and blood coagulation. BTL.010 DNA is useful in gene therapy. The present sequence is a BTL.010 protein related protein.
                                                                                                                      diagnostic
                                                                                                                                                sarcomatoid
                                                                                                                                                                           cytostatic; gene therapy; differential expression; renal cell carcinoma; clear cell RCC; papillary RCC; chromophobe/oncocytoma RCC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    glomerulonephritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          idiopathic pulmonary fibrosis; adult respiratory distress syndrome; cystic fibrosis; rheumatoid arthritis; organ failure; glomerulonephritis; inflammatory disease; oxidative burst; platelet activation;
        WO2004032842-A2
                                                                                                                                                                                                                                                                                                                         15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                 ADOS5146;
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADO55146 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 15; 17pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated polynucleotide encoding BTL.010 serine proteinase, usefu for treating e.g. emphysema, idiopathic pulmonary fibrosis, adult respiratory distress syndrome, cystic fibrosis, rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (FARB ) BAYER PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coagulation; gene therapy.
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                                                                                                                                                                                                                                                                 #48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                      RCC;
                                                                                                                                                                                                                                                                 with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            $
                                                                                                                                                                                                                                                                                                                         (first
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                                                                                                                                                   papillary RC; Wilms'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ū
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                                                                                                                                                                                                                                                           increased gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note
                                                                                                                                                                                                                                                                                                                                                                                                                                      protein;
                                                                                                                                                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roczniak S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         80.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      = Kunitz domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 264;
Pred. No. 2
                                                                                                                                                tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                      A
                                                                                                                                             gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 8;
2.1e-23;
                                                                                                                                                expression; kidney cancer
                                                                                                                                                                                                                                                              in renal cell carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 189
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                                                            Query Match
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Matches 48
                                                                                                                                                   The invention relates to novel genes that are differentially expressed i sub-types of renal cell carcinomas and methods of detecting them using nucleic acids and probes. The nucleic acid probes hybridize with part or all of a coding sequence that is overexpressed in clear cell renal cell carcinoma (CC-RCC) papillary RCC, chromophobe/oncocytoma RCC, excomatoid RCC, TCC, or Wilms' tumors, which overexpression is based on comparison to a baseline value. The methods and compositions of the present invention are useful in the field of molecular biology and medicine, in particular for gene expression profiling of certain types okidney cancer, in identifying diagnostic markers, and treating such cancer patients. This sequence corresponds to the protein encoded by a gene with increased expression in CC-RCC.
                                                                                                                            Sequence 1235 AA;
                                                                                                                                                                                                                                                                                                                                                                        Example IV; SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                  New nucleic acid and polypeptide compositions, useful in the field molecular biology and medicine, in particular for gene expression profiling, identifying diagnostic markers, and treating certain ty
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-OCT-2002; 2002US-0415775P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-OCT-2003; 2003WO-US031476
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (VAND-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2004-340789/31.
                                                              48;
                  1 YPVRCILPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VAN
                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADO54960.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Takahashi M;
YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANDEL INST
                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                           80.2%;
                                                                                                                                                                                                                                                                                                                                                                        243;
                                                                                                                                                                                                                                                                                                                                                                     53pp;
                                                              <u>,</u>
                                                            Score 264; DB 8;
Pred. No. 1.4e-22;
0; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                       English
                                                              10;
                                                                                          Length 1235;
                                                               Indels
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RESULT 14
ABR58408
ID ABR58
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EM Huma
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EM
     02-OCT-2001;
05-OCT-2001;
09-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; NOV; antidiabetic; anorectic; antibacterial; virucide; immunomodulator; cytostatic; nootropic; neuroprotective; dyslipidaemia; antiparkinsonian; antilipaemic; gene therapy; metabolic disorder; diabetes; obesity; infection; cachexia; cancer; Parkinson's disease; neurodegenerative disorder; Alzheimer's disease; immune disorder;
                                                                                                                                                                                                                                                                                                              10-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human NOV22a.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JUL-2003
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     2001US-0326483P.
2001US-0327342P.
2001US-0327917P.
                                                                                                                                                                                                            2002WO-US031358
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RESULT 15
ADB60631
ID AB60
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09-OCT-2001;
19-OCT-2001;
12-OCT-2001;
15-OCT-2001;
17-OCT-2001;
22-OCT-2001;
24-OCT-2001;
24-OCT-2001;
24-OCT-2001;
10-NOV-2001;
11-APR-2002;
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11-APR-2002;
11-APR-2002;
                                                                                                                                                                            27-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to novel human NOV proteins and their coding sequences (ACC72075-ACC72181 and ABRS363) ABRS366). The NOV proteins are useful in manufacturing a medicament for treating a syndrome associated with a human disease. The NOV proteins and coding sequences may be used to diagnose, treat or prevent metabolic disorders such as diabetes or obesity, infections, cachexia, cancer, neurodegenerative disorders such as Alzheimer's disease or Parkinson's disease, immune disorders, haematopoietic disorders and various dyslipidaemias
  Human BTL.010; neutral serine protease inhibitor; elastase inhibitor; proteinase-3 inhibitor; Kunitz domain; emphysema; idiopathic pulmonary fibrosis; adult respiratory distress syndrome; cystic fibrosis; rheumatoid arthritis; organ fallure; glomerulonephri
                                                                                                                                                                                                                             AAB60631;
                                                                                                                                                                                                                                                                           AAB60631 standard; protein; 51 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NOVX polypeptides and nucleic acids useful for diagnosing, preventing treating NOVX-associated disorders, e.g. diabetes, obesity, cancer or dyslipidemia, and in chromosome mapping, tissue typing or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Alsobrook JP, Anderson K, Ellerman K,
                                                                                                                           Human protease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1280 AA;
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16-MAY-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vernet
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                                                                                                                                                                                                                                                                                                                                                                                                  752 YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 809
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         48;
                                                                                                                                                                                                                                                                                                                                                                                                                                      1 YPVRCILIPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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M VOSS EZ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; 2001US-0328029P.
2001US-0328044P.
2001US-0328849P.
2001US-032914P.
2001US-0330142P.
2001US-0341058P.
2001US-0349575P.
2001US-0349575P.
2002US-0371980P.
2002US-0371980P.
2002US-0371935P.
2002US-0371935P.
2002US-0371935P.
2002US-0371935P.
2002US-0371935P.
2002US-0371935P.
2002US-0371935P.
2002US-0371935P.
2002US-0371935P.
2002US-0381101P.
2002US-038339P.
2002US-038339P.
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                                                                                                                                                                          (first entry)
                                                                                                                         inhibitor BTL.010 Kunitz domain fragment,
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K, Gerlach VL, Gorn
Li L, Miller CE, P
ts RA, Smithson G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zerhusen BD,
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Pred. No. 1.4e-22;
0; Mismatches 10
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VL, Gorman L, Guo X,
r CE, Patturajan M, R
hson G, Spytek KA, Ta
, Zhong M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10;
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glomerulonephritis;
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Taupier
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                                                                                                                                                            database using the Kunitz domain sequences AAB60630, and was confirmed as being novel using the Kunitz domain sequences AAB60632, and AAB60633.

This sequence information was extended to provide a larger region of BTL-010 protein sequence data (AAB60634) by identifying an open reading frame (ORF) which comprised DNA encoding the BTL-010 Kunitz domain corresponding to bases 16016-16414 of GenBank accession number AC004846.

The entire BTL-010 Kunitz domain sequence (AAB60623) was obtained from the BTL-010 ORF-encoded sequence. The BTL-010 protein, and pharmaceutical compositions comprising it, may be used for inhibiting protease activity, particularly that of leukocyte elastase, in the prevention, treatment or amelioration of medical conditions such as emphysema, idiopathic pulmonary fibrosis, adult respiratory distress syndrome, cystic fibrosis, rheumatoid arthritis, organ failure or glomerulonephritis. BTL-010 condition such as platelet activation, blood coagulation, neutrophil activation, or monocyte activation, blood coagulation, neutrophil activation, or monocyte activation, blood coagulation, neutrophil activation of the treatment of inflammatory diseases and diseases involving lung and vascular injury. The present sequence represents a substantial proportion of the human BTL-010 protease inhibitor Kunitz domain
                                                                                Matches
                                                                                                   Query Match
Best Local :
                                                                                                                                                            Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       platelet activation; blood coagulation; neutrophil activation; monocyte activation; angioplasty; inflammatory disease; lung injury; vascular injury; nephrotropic; antirheumatic; antiarthritic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neutral serine proteases such as elastase and proteinase-3, relative to trypsin-like and chymotrypisn-like proteases. A substantial proportion of the BTL.010 protein Kunitz domain (AAB60631) was identified via homology searching in the GenBank high throughput genomic (HTG) DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel serine proteinase inhibitor of the Kunitz family, BTL.010 useful for treating emphysema, cystic fibrosis, adult respiratory distress
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB60634).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Col 5-6; 17pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for treating emphysema, cystic fibrosis, adult respiratory distress syndrome, rheumatoid arthritis, organ failure and glomerulonephritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB; AAF59750
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Davies C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DAVI/) DAVIES C. (CHEN/) CHEN D. (ROCZ/) ROCZNIAK S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to
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                                                                                42;
                                           ഗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            family,
                                                                                                   Similarity
                          CLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQ 56
CLLPSAHGSCADWAARWYFVASVGQCNRFWYGGC-GNANNFASEQECMSSCQ 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ily, BTL.010 (fragments given in AAB60623, AAB60631 and The BTL.010 protein is thought to preferentially inhibit
                                                                                                                                                            ₽,
                                                                              Conservative
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                                                                                                 67.0%;
80.8%;
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                                                                              <u>.</u>
                                                                                                 Score 220.5;
Pred. No. 8.7
                                                                                Mismatches
                                                                        . 8.7e-19;
9;
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n in AAB60623, AAB60631
                                                                                                                   Length 51;
                                                                              Indels
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                                                                              Gaps
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Search Job tim completed: August 25, 2005, 04:52:21

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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                          Database :
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Maximum DB seq length: 2000000000
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Perfect score:
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PIR_79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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329
                                                                                                                                                                                                                                                                      283416 seqs, 96216763 residues
                                                                                                                                                                                                                                                                                                        BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                   August 25, 2005, 04:48:01; Search time 16 Seconds (without alignments) 348.786 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                1 YPVRCLLPPATGPCKARIIR......RGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                       283416
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	BG	ID	Description
۲ !	192	58.4	100	<u>- </u>	TIBO	basic proteinase i
N	190	57.8	59	N	S00371	rotinin G1 -
ω	188	57.1	58	N	S10063	2
4	184	55.9	60	H	TIBOR	serum basic protei
σ	184	٠	100	۳	TIBOSP	spleen basic prote
თ	174		122	_	A55115	uterine plasmin/tr
7	168	•	111	N	S41082	amyloid precursor
æ	168	٠	751	N	A49974	beta-amyloid precu
9	168	•	763	N	A49321	amyloid beta (A4)
10	168	٠	765	N	S42880	amyloid precursor-
11	164	•	61	<u>ب</u>	TIVIT1	asic prote
12	163	٠	57	N	A59204	basic proteinase i
14	163	49.5	110	۸ د	A44180	talcatoxin serine
15	161		249	N	T32060	etical prot
16	159		61	_	TIHCBP	proteinase inhibit
17	159	•	62	N	S07451	proteinase inhibit
18	159		252	N	JG0185	
19	158	•	82	_	TIHABK	å
20	158	48.0	76	N	S06678	128
21	158	•	76	N	S03607	Alzheimer's diseas
22	158	٠	484	4	A32761	hypothetical Alzhe
23	158	٠	770	هــو	QRHUA4	Alzheimer's diseas
24	157	•	76	N	S04855	Alzheimer's diseas
25	157		100	N	A32282	œ
26	156		2225	N	T26063	hypothetical prote
27	155	٠	747	N	JH0773	Alzheimer's diseas
28	154	46.8	123	N	A29652	a-tryps
29	152	46.2	265	N	A53390	Kunitz-type protei

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	
144	144	145	145	147	148	148	148	148	148	150	151	151	151	151	152	
43.8	43.8	44.1	44.1	44.7	45.0	45.0	45.0	45.0	45.0	45.6	45.9	45.9	45.9	45.9	46.2	
302	57	922	352	304	2167	1558	235	65	64	67	352	349	125	62	349	
<u>, , , , , , , , , , , , , , , , , , , </u>	۲	N	1	بر	N	N	N	r	N	Н	,	N	μ	N	N	
TIRTGK	TINJVC	T23573	HCHU	TIHUGK	T34395	C89114	A54951	TIVIVC	S41399	TIBOC	TIBOBI	S35708	TIHOBI	S19327	S21089	
tissue factor path	venom basic protei	hypothetical prote	alpha-1-microglobu	tissue factor path	hypothetical prote	protein C37C3.6a {	tissue factor path	venom basic protei	Kunitz-type protei	trypsin inhibitor,	alpha-1-microglobu	alpha-1-microglobu	alpha-1-microglobu	venom basic protei	_alpha-1-microglobu	

ALIGNMENTS

A; Molecule type: protein A; Residues: 36-93 < IXE> R; Rassell, B; Laskowski, M. Biochem. Biophys. Res. Commun. 20, 463-468, 1965 A; Title: The basic trypsin inhibitor of bovine pancreas. V. The disulfide linkages. A; Reference number: A90162; MUID:66083012; PMID:5860161 A; Contents: annotation; disulfide bonds A; Accession: A90162 A; Molecule type: protein A; Residues: 36-93 < KAS>	A; Molecule type: protein A; Residues: 36-93 <fio> R; Residues: 36-93 <fio> R; Kekita, M.; Jone, C.S.; Kamo, M.; Tsugita, A.; Kizuki, K.; Moriya, H. Protein Seq. Data Anal. 5, 7-11, 1992 A; Title: Purification and characterization of the major cationic kallikrein inhibitor in A; Reference number: S28197; MUID:93150003; PMID:1283464 A; Accession: S28197</fio></fio>	A;Accession: S10546 A;Molecule type: DNA A;Mesidues: 34-97 <kin> A;Fesidues: 34-97 <kin> R;Floretti, E.; Angeletti, M.; Florucci, L.; Barra, D.; Bossa, F.; Ascoli, F. R;Floretti, E.; Angeletti, M.; Florucci, L.; Barra, D.; Bossa, F.; Ascoli, F. Biol. Chem. Hoppe-Seyler 369(Suppl.), 37-42, 1988 A;Title: Aprotinin-like isoinhibitors in bovine organs. A;Reference number: S02485; MUID:89076531; PMID:2462435 A;Accession: S02486</kin></kin>	A; Molecule type: DNA A; Molecule type: DNA A; Residues: 1-100 < CRE> A; Cross-references: GB:M20934; GB:X05274; NID:g162767; PIDN:AAD13685.1; PID:g162769 A; Cross-references: GB:M20934; GB:X05274; NID:g162767; PIDN:AAD13685.1; PID:g162769 A; Kingston, I.B.; Anderson, S. Biochem. J. 233, 443-450, 1986 Biochem. J. 233, 443-450, 1986 A; Title: Sequences encoding two trypsin inhibitors occur in strikingly similar genomic er A; Reference number: S10546; MUID:86158754; PMID:2420326	A; Molecule type: DNA; mRNA A; Residues: 1-100 < CR2> A; Cross-references: UNIPROT: P00974; GB:M20934; GB:X05274; NID:g162767; PIDN:AAD13685.1; I R; Creighton, T.B.; Charles, I.G. Cold Spring Harb. Symp. Quant. Biol. 52, 511-519, 1987 A; Title: Biosynthesis, processing, and evolution of bovine pancreatic trypsin inhibitor. A; Reference number: A90926; MUID:88295740; PMID:2456884 A. Accession. A20333	RESULT 1 TIBO TIBO basic proteinase inhibitor precursor - bovine basic proteinase inhibitor precursor - bovine N;Alternate names: aprotinin; basic pancreatic trypsin inhibitor; BPTI; cationic kallikre C;Species: Bos primigenius taurus (cattle) C;Date: 24-Apr-1984 #sequence revision 22-Jul-1994 #text change 09-Jul-2004 C;Accession: \$00277; A30333; \$10546; \$02486; \$28197; A90162; A92023; A90736; A90927; A346 R;Creighton, T.E.; Charles, I.G. J. Mol. Biol. 194, 11-22, 1987 A;Title: Sequences of the genes and polypeptide precursors for two bovine protease inhibi A;Reference number: \$00274; MUID:87283904; PMID:2441071 A;Accession: \$00274

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R;Anderer, F.A.; Hornle, S.
J. Biol. Chem. 241, 1568-1572, 1966
A;Title: The disulfide linkages in kallikrein inactivator of bovine lung.
A;Reference number: A9203; MUID:66171231; PMID:5296424
A;Contents: annotation; disulfide bonds
A;Accession: A92023
A;Molecule type: protein
A;Residues: 36-93 <AN2>
R;Chauvet, J.; Acher, R.
Bull. Soc. Chim. Biol. 49, 985-1000, 1967
A;Title: La structure covalente d'un inhibiteur polypeptidique de la trypsine (inhibiteu A;Reference number: A90736; MUID:68012003; PMID:6053284
A;Contents: annotation; disulfide bonds
A;Accession: A90736
A;Molecule type: protein
A;Residues: 36-93 <CHA>
A;Residues: 36-93 <CHA>
A;Residues: 36-93 <CHA>
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A;Molecule type: DNA
A;Residues: 'PSLFNRDPPIPA', 34-97, 'GKTGGRAEGEGKG' <AND>
A;Residues: 'PSLFNRDPPIPA', 34-97, 'GKTGGRAEGEGKG' <AND>
A;Cross-references: GB.X03365; GB.K00966; NID:g142; PIDN:CAA27062.1; PID:g1364183
R;Siekmann, J.; Wenzel, H.R.; Schroeder, W.; Tschesche, H.
Biol. Chem. Hoppe-Seyler 369, 157-163, 1988
Biol. Chem. Hoppe-Seyler 369, 157-163, 1988
A;Title: Characterization and sequence determination of six aprotinin homologues from A;Reference number: S00371; MUID:88221840; PMID:2453200
A;Accession: S10062
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A;Residues: 36-53,55-81 <LEW>
A;Residues: 36-53,55-81 <LEW>
R;Anderson, S.; Kingston, I.B.
Proc. Natl. Acad. Sci. U.S.A. 80, 6838-6842, 1983
Prot. Natl. Acad. Sci. U.S.A. 80, 6838-6842, 1983
A;Title: Isolation of a genomic clone for bovine pancreatic trypsin inhibitor by A;Reference number: A93977; MUID:84070725; PMID:6580617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: protein
A;Residues: 36-66,'p',68-82,'S',84-93 <SIE>
A;Experimental source: lung
A;Note: the authors designated this protein as isoaprotinin 2
C;Comment: Basic proteinase inhibitor is an intracellular polypeptide found
C;Genetics:
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A;Contents: annotation; X-ray crystallography of basic protease inhibitor, 2.5 angstroms R;Lewis, R.V.; Ray, P.; Coguill, R.; Kruggel, W.
Biochem. Biophys. Res. Commun. 167, 543-547, 1990
A;Title: Presence of pancreatic trypsin inhibitor in adrenal medullary chromaffin cells. A;Reference number: A34658; MUID:90211226; PMID:2322242
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A;Residues: 36-93 <DLO>
R;Huber, R; Kukla, D; Ruhlmann, A.;
Naturwissenschaften 57, 389-392, 1970
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Accession: A93977
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                                                                                                                                                                                                                                                                                                                                                                                      Superfamily: basic proteinase inhibitor; animal Kunitz-type prot

;Keywords: serine proteinase inhibitor;

;1-20/Domain: signal sequence #status predicted <SIG>

;21-35/Domain: propeptide #status predicted <PRO>

;21-35/Domain: propeptide #status predicted <PRO>

;36-10/Product: basic proteinase inhibitor #status experimental

;40-90/Domain: animal Kunitz-type proteinase inhibitor homology <

;40-90/49-73,65-86/Disulfide bonds: #status experimental
                                                                                                                                                                                                                                                                                                                                                         :50/Inhibitory site: Lys (trypsin, chymotrypsin, kallikrein, plasmin)
                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                       Local Similarity
                                                                               37
                                                                                                                                                                                                                          33;
                                                                                                                                                     N
                                                                                                                       PVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                            PDFCLEPPYTGPCKARIIRYFYNAKAGLCOTFVYGGCRAKRNNFKSAEDCMRTCGGA
                                                                                                                                                                                                                          Conservative
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Pred. No. 2
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.5e-16;
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FEBS Lett. 119, 58-62, 1980
A; Title: A new Kunitz-type:
A; Reference number: A01206;
                                                                                                                                                                                                                                                                                                                                                                                          serum basic proteinase inhibitor - bovine C;Species: Bos primigenius taurus (cattle) C;Date: 31-Oct-1980 #sequence_revision 31-
A;Cross-references: UNIPROT:P00975
C;Comment: This inhibitor has acti
C;Superfamily: basic proteinase in
                                                                                                                                                                                                                                                                                                                                                         C; Accession: A01206
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A;Molecule type: protein A;Residues: 1-60 <WAC>

A;Accession: A01206

#sequence_revision 31-Oct-1980 #text_change 09-Jul-2004

Hochstrasser, K.; Lempart, K.; Geiger,

inhibitor from bovine serum

MUID:81044408;

PMID:7428928

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determination

activity very se inhibitor; a

similar to that of the animal Kunitz-type prote

of the basic proteinase

protease inhibitor

inhibit br homolo

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isoaprotinin G2 - bov
C,Species: Bos primig
C,Date: 31-Dec-1990 #
C,Accession: S10063
R,Siekmann, J.; Wenze
RESULT
TIBOR
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S10063
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C;Superfamily: basic proteinase inhibitor; animal Kunitz-type proteinase C;Superfamily: basic proteinase inhibitor; animal kunitz-type proteinase inhibitor c;Keywords: pyroglutamic acid; serine proteinase inhibitor homology <BPI>F;6-56/Domain: animal Kunitz-type proteinase inhibitor homology <BPI>F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status predicted F;16/Inhibitory site: Lys (trypsin) #status predicted
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A;Title: Characterization and sequence determination of A;Reference number: S00371; MUID:88221840; PMID:2453200 A;Accession: S00371
A;Molecule type: protein A;Residues: 1-59 <SIE>
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C;Species: Bos primigenius indicus x Bos primigenius taurus (cattle)
C;Date: 31-Dec-1990 #sequence_revision 31-Dec-1990 #text_change 09-J-C;Accession: S00371
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S00371
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                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: protein A; Residues: 1-58 < SIE>
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                                                                                                  PDFCLEPPYTGPCKARMIRYFYNAKAGLCQPFVYGGCRAKRNNFKSAEDCMRTCGGA
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primigenius indicus x Bos primigenius taurus (cat
primigenius indicus x Bos primigenius taurus (cat
primigenius taurus (cat
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56.1%;
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Pred. No. 4.7e-16;
8; Mismatches 17
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Pred. No. 2.7e-16;
9; Mismatches 16
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A; Kreshandes: 1-Du KCKA2
R(Cresighton, T.B.; Charles, I.G.
R(Classing Harb, Symp. Quant. Biol. 52, 511-519, 1987)
A; Title: Blosynthesis, processing, and evolution of bovine pancreatic trypsin inhibitor.
A; Reference (Biological Muld: 1925/40; PMID:2456884
A; Recession: B3033)
A; Molecula type: Dal
A; Residues: 1-100 KCRR>
A; Residues: 1-100 KCRR>
A; Residues: 1-100 KCRR>
A; Title: Primary structure of a procease isoinhibitor from bovine spleen. A possible int A; Reference number: 127417; MUID:88007630; PMID:3554647
A; Recession: R127417
A; RAccession: R127417
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A; RAccession: R127417
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C;Species: Bos primigenius taurus (cattle)
C;Date: 07-Sep-190 #sequence revision 22-Jul-1994 #text change 09-Jul-2004
C;Accession: S00274; B30333; Ā27417; S02487; S10547; S13478; A23915; S10064
R;Creighton, T.E.; Charles, I.G.
J. Mol. Biol. 194, 11-22, 1987
A;Title: Sequences of the genes and polypeptide precursors for two bovine planetering number: S00274; MUID:87283904; PMID:2441071
A;Accession: S00274
A;Molecule type: DNA; mRNA
A;Residues: 1-100 <CR2>
A;Residues: 1-10
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Pred. No. 1.5e-15;
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                                                                  A;Title: Expression, purification and characterization of a A;Reference number: S41082; MUID:94139895; PMID:8307156 A;Accession: S41082
                                                                                                                                                                                                                 C;Accession: S41082
R;Petersen, L.C.; Bjorn
FEBS Lett. 338, 53-57,
                                                                                                                                                                                                                                                                                                                                                                 C;Species: Homo sapiens (man)
C;Date: 25-Dec-1994 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C; Keywords: seri
F;38-88/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Note: authors translated the codon GGC for residue 36 as Ala, AGC for r C;Superfamily: basic proteinase inhibitor; animal Kunitz-type proteinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:Q29100; GB:L14282; NID:g682652; PIDN:AAA62425.1; PID:g682653
A;Note: authors translated the codon GGC for residue 36 as Ala, AGC for residue 48 as Ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amyloid precursor protein homolog -
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35

Bjorn,

S.E.;

Norris,

Έ :

Κ.,

Kunitz-type protease inhibito

Foster,

D.C.

03-Aug-1995 #text_change

09-Jul-2004 ი.;

human

(fragment)

N 28;

Conservative

8;

Score 174; DB 1; Pred. No. 4.8e-14; 8; Mismatches 18

18;

Gaps

0

Length 122;

52.9**%**; 51.9**%**;

PVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSC 55

PGFCREPPYTGPCSAHFVRYFYNATTGLCQSFVYGGCRGKQNNFMDEKECLHTC

serine proteinase inhibitor min: animal Kunitz-type prote

proteinase inhibitor homology

<BPI>

inhibitor homolo

Arc

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A;Experimental source: lung C;Comment: This inhibitor is found in many bovine organs. In spleen, it is located in vat C;Comment: Compared with the basic proteinase inhibitor, affinity for trypsin and chymoti C;Comment: Inhibitors I and III are intermediates in the processing of the precursor to t C;Genetics:
                                                                                                                                                                             R;Stallings-Mann,
J. Biol. Chem. 26
                                                                                                                                                                                                                                    uterine plasmin/trypsin inhibitor - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 09-Jul-2004
C;Accession: A55115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             F;34-98/Product: spleen inhibitor I #status experimental <SI1> F;36-97/Product: spleen inhibitor III #status experimental <SI3> F;36-97/Product: spleen inhibitor II #status experimental <SI2> F;40-90/Domain: animal Kunitz-type proteinase inhibitor homology <BPI> F;40-90,49-73,65-86/Disulfide bonds: #status predicted F;50/Inhibitory site: Lys (trypsin, chymotrypsin, kallikrein) #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Biol. Chem. Hoppe-Seyler 369, 157-163, 1988
A;Title: Characterization and sequence determination of six aprotinin homologues from bov A;Reference number: S00371; MUID:88221840; PMID:2453200
                                                                                                                A;Title: Purification, characterization, and cDNA cloning of a Kunitz-type proteinase in
A;Reference number: A55115; MUID:95014140; PMID:7929061
                                                                                                                                                  J. Biol. Chem. 269, 24
A; Title: Purification,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        F;1-20/Domain: signal sequence #status predicted < F;21-33/Domain: propeptide #status predicted <PRO>
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C; Superfami
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: protein A; Residues: 36-93 <SIE>
A; Residues: 1-122 <STA>
                           A; Molecule type: mRNA
                                                         A; Status: preliminary
                                                                                           A; Accession: A55115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Keywords:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Accession: S10064
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PVRCILPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PDFCLEPPYTGPCKAKMIRYFYNAKAGFCETFVYGGCKAKSNNFRSAEDCMRTCGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                basic proteinase inhibitor; animal Kunitz-type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                               M.L.; Burke, M.G.;
                                                                                                                                                                             24090-24094, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        52.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 184; DB 1;
Pred. No. 2.4e-15;
                                                                                                                                                                                                            Trout, W.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                               Roberts,
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A;Cross-references: EMBL:Z22572; NID:g394763; PIDN:CAA80295.1; PID:g394764 R;Wasco, W.; Gurubhagavatula, S.; Paradis, M.; Romano, D.M.; Sisodia, S.S. Nature Genet. 5, 95-99, 1993 A;Title: Isolation and Characterization of APLP2 encoding a homologue of tipA;Reference number: S40519; MUID:94035131; PMID:8220435
                                                                                                                                                                                                                                                                  A;Note: sequence extracted from NCBI backbone (NCBIN:131198, NA;Note: expression was shown in placenta, brain, heart, lung, R;von der Kammer, H.; Klaudiny, J.; Hannes, J.; Scheit, K.H. submitted to the EMBL Data Library, April 1993
A;Description: The human homologue of the murine CDEI binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              beta-amyloid precursor protein 2 homolog APLP2 - mouse C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 06-Oct-1994 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Accession: A49974
R;Slunt, H.H.; Thinakaran, G.; Von Koch, C.; Lo, A.C.; Tanzi, R.E.; Sisodia, S.S.
J. Biol. Chem. 269, 2637-2644, 1994
A;Title: Expression of a ubiquitous, cross-reactive homologue of the mouse beta-amyloid A;Reference number: A49974; MUID:94132029; PMID:8300594
A;Accession: A49974
A;Accession: A49974
A;Accession: A49974
A;Status: preliminary; not compared with conceptual translation
A;Residues: 1-751 <SLU>
A;Cross-references: UNIPROT:Q60709; GB:U15571; NID:g558467; PIDN:AAA50603.1; PID:g558468
                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: mRNA
A;Residues: 1-763 <SPR>
A;Residues: 1-763 <SPR>
A;Cross-references: UNIPROT:006481; GB:S60099; NID:g300168; PIDN:AAC60589.1; PID:g300169
A;Experimental source: placenta
                                                                                                                                                     A; Molecule type: mRNA
A; Residues: 1-763 < VON>
                                                                                                                                                                                                                A;Reference number: S34644
A;Accession: S34644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Reference number: A49321;
A; Accession: A49321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G;Date: 24-Feb-1994 #sequence revision 18-Nov-1994 #text_change C;Accession: A49321; S34644; S40519 R;Sprecher, C.A.; Grant, F.J.; Grimm, G.; O'Hara, P.J.; Norris, Biochemistry 32, 4481-4486, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: UNIPROT:060709; GB:U15571; NID:g558467; PIDN:AAA50603.1; PID:g558468
A;Note: sequence extracted from NCBI backbone (NCBIP:144636)
C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase i
F;310-360/Domain: animal Kunitz-type proteinase inhibitor homology <BPI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       밁
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A;Residues: 1-111 <PET>
A;Cross-references: UNIPROT:Q7M4L3
F;59-109/Domain: animal Kunitz-type proteinase inhibitor homology <BPI>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;Title: Molecular cloning of the cDNA for a human amyloid precursor;Reference number: A49321; MUID:93250009; PMID:8485127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CSQEAMTGPCRAVMPRWYFDLSKGKCVRFIYGGCGGNRNNFESEDYCMAVCK 110
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No. 1.4e-12;
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6; Mismatches
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Pred. No. 2.4e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                        protein is an amyloid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            F.; Norris,
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                                  of the Alzheimer'
                                                                                          S.S.;
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                                                                                       Hyman,
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                                                                                             B.T.
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A;Description: Complete nucl
A;Reference number: 942880
A;Reference number: 942880
A;Accession: 942880
A;Molecule type: mRNA
A;Residues: 1-765 <SAN>
                                                                                                                                                   A;Cross-references: EMBL:X77934
C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase C;Keywords: alternative splicing
F;312-362/Domain: animal Kunitz-type proteinase inhibitor homology <BPI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Rattus norvegicus (Norway rat)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1
C;Accession: S42880; S47528
R;Sandbrink, R; Masters, C.L.; Beyreuther, K. submitted to the EMBL Data Library, March 1994
A;Description: Complete nucloetide ad deduced &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Gene: GDB:APLP2; APPL2
A;Gene: GDB:APLP2; APPL2
A;Cross-references: GDB:139159; OMIM:104776
A;Map position: 11q23-11q25
C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type C;Keywords: alternative splicing; transmembrane protein C;Keywords: animal Kunitz-type proteinase inhibitor homology <BPI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Accession: $40519
A;Status: preliminary
A;Molecule type: mkNA
A;Molecule type: mkNA
A;Residues: 1-763
A;Cross-references: GB:L27631; NID:g450391; PIDN:AAC41701.1; PID:g450392
C;Genetics: 1-7631; NID:g450391; PIDN:AAC41701.1; PID:g450392
밁
                                                                                                                                                                                                                                                                                                 RESULT
S42880
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                                                                                                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 1-765 <SA2>
                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:P15943; EMBL:X77934 R;Sandbrink, R.; Masters, C.L.; Beyreuther, K.
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Best Local &
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312
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                                                                                                 Similarity
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                      CLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQ 56
CSQEAMTGPCRAVMPRWYFDLSKGKCVRFIYGGCGGNRNNFESEDYCMAVCK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CSQEAMTGPCRAVMPRWYFDLSKGKCVRFIYGGCGGNRNNFESEDYCMAVCK 361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQ 56
                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                           51.1%;
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                                                                           Score 168; DB 2;
Pred. No. 1.4e-12;
6; Mismatches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 168;
Pred. No.
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venom basic proteinase inhibitor I - western sand viper N;Alternate names; venom trypsin inhibitor I C;Species: Vipera ammodytes ammodytes (western sand viper) C;Date: 19-Feb-1984 #sequence_revision 19-Feb-1984 #text_change C;Accession: A01222

A;Title: The primary structure of Vipera ammodytes venom A;Reference number: A01222; MUID:84053385; PMID:6639951 A;Accession: A01222

R;Ritonja, A.; Meloun, B.; Gubensek, Biochim. Biophys. Acta 748, 429-435,

1983

trypsin

inhibitor

09-Jul-2004

ij

A; Residues: 1-61 <R A; Cross-references: A; Molecule type: protein A; Residues: 1-61 < RIT>

UNIPROT: P00991

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submitted to the Protein Sequence Database, March 2000
A, Description: Lymnaea trypsin inhibitor.
A; Reference number: A59204
A; Accession: A59204
A; Accession: A59204
A; Molecule type: protein
A; Residues: 1-57 < NAG>
A; Cross-references: UNIPROT:07M411
A; Experimental source: albumen gland
C; Superfamily: basic proteinses inhibitor; animal Kunitz-!
C; Reywords: 91ycoprotein; serine proteinase inhibitor
F; 5-55/Domain: animal Kunitz-type proteinase inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N;Alternate names: trypsin inhibitor - great pond snail
C;Species: Lymnaea stagnalis (great pond snail)
C;Date: 04-Mar-2000 #sequence_revision 04-Mar-2000 #text_change 09-Jul-2004
C;Accession: A59204
R;Nagle, G.
                                                                                                                                                                                                                                       R;Possani, L.D.; Martin, B.M.; Yatani, A.; Mochca-Morales, J.; Zamudio, F.Z.; Gurrola, G. Toxicon 30, 1343-1364, 1992
A;Title: Isolation and physiological characterization of taicatoxin, a complex toxin with A;Reference number: A44180; MUID:93134601; PMID:1485334
A;Accession: A44180
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-62 <POS>
A;Cross-references: UNIPROT:Q7LZE4
                                                                                                 A;Experimental source: subsp. scutellatus, venom R;Experimental source: subsp. scutellatus, venom R;Experimental source: sequence extracted from NCBI backbone (NCBIP:122482) C;Superfamily: basic proteinase inhibitor; animal Kunitz-type proteinase C;Keywords: serine proteinase inhibitor c;F;7-57/Domain: animal Kunitz-type proteinase inhibitor homology <BPI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               taicatoxin serine proteinase inhibitor component - Australian taipan C;Species: Oxyuranus scutellatus scutellatus (Australian taipan) C;Date: 27-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004 C;Accession: A44180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Comment: This protein inhibits trypsin and kallikrein.
C;Superfamily: basic proteinase inhibitor; animal Kunitz-type proteinase inhibitor homol
C;Keywords: pyroglutamic acid; serine proteinase inhibitor; venom
F;7-57/Domain: animal Kunitz-type proteinase inhibitor homology <BPI>
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;7-57,16-40,32-53/Disulfide bonds: #status predicted
F;17/Inhibitory site: Lys (trypsin) #status predicted
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Query Match
Best Local S
Matches 28
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  Similarity 28; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSC
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  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10;
7;
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                         Score 163; DB 2;
Pred. No. 5.6e-13;
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     Mismatches
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A;Cross references: UNIPROT:P00993
C;Comment: This inhibitor, isolated from egg white, consists of two nonhomologous domains C;Superfamily: loggerhead basic proteinase inhibitor; animal Kunitz-type proteinase inhibitor
C;Keywords: pyroglutamic acid; serine proteinase inhibitor
F;8-58/Domain: animal Kunitz-type proteinase inhibitor homology <BPI>
F;63-105/Domain: antileukoproteinase repeat homology <ALP>
F;11/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;8-58,17-41,33-54,67-93,76-97,80-92,86-101/Disulfide bonds: #status predicted
F;18/Inhibitory site: Lys (trypsin) #status predicted
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                                                                                                                                                                              A;Gene: CESP:R12A1.3
A;Map position: 5
A;Introns: 75/1; 139/1
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                                                                                                                                                                                                                                                                                                                                                            A;Status: preliminary; translated from
                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Pauley, A.; Andrews, S. submitted to the EMBL Data Library, July : A;Description: The sequence of C. elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypothetical protein R12A1.3 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #tC;Accession: T32060
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C;Accession: A01224
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                                                                                                                                                                                                                                                                                                                                              A; Molecule type: DNA
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C;Species: Caretta caretta (loggerhead)
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143
                          5 CLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS
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CSLPLAVGSCTAPAVRFYYDASSGRCNQFMYSGCGGNANNFQS---
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                                                                                        Conservative
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                                                                                                             48.98;
55.68;
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                                                                                                             Score 161; DB 2;
Pred. No. 3.5e-12;
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Search completed: August Job time : 17 secs

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2005, 04:55:46

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Result
No.
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Maximum
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence:
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Perfect score:
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length: 2000000000
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173.186 Million cell updates/sec
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329
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60.2
60.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/BCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/backfiles1.pep:*
                                                     GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BB
                                                 US-09-369-494-1
US-09-369-494-15
US-09-369-670-15
US-09-369-670-15
US-09-369-670-18-75
US-08-084-718-75
US-08-084-718-75
US-08-084-718-79
US-08-443-977-61
US-08-443-977-61
US-08-443-977-61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      513545
Sequence 75, Appl
Sequence 75, Appl
Sequence 59, Appl
Sequence 59, Appl
Sequence 59, Appl
Sequence 71, Appl
Sequence 71, Appl
Sequence 71, Appl
Sequence 69, Appl
Sequence 69, Appl
Sequence 41, Appl
Sequence 41, Appl
Sequence 41, Appl
Sequence 61, Appl
Sequence 77, Appl
Sequence 77, Appl
Sequence 81, Appl
Sequence 77, Appl
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Sequence 1, Appli
Sequence 15, Appl
Sequence 15, Appl
Sequence 9, Appli
Sequence 9, Appli
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	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28
	192	192	192	192	193	193	193	193	193	193	193	193	193	193	193	194	194	194
	58.4	58.4	58.4	58.4	58.7	58.7	58.7	58.7	58.7	58.7	58.7	58.7	58.7	58.7	58.7	59.0	59.0	59.0
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ALIGNMENTS	US-08-084-718-39	US-08-084-718-1	US-07-598-737C-1	US-07-664-989B-44	US-08-443-977-73	US-08-443-977-65	US-08-443-976-73	US-08-443-976-65	US-08-084-718-73	US-08-084-718-65	US-08-896-322-5	US-08-443-977-34	US-08-443-976-34	US-08-084-718-34	US-07-598-737C-33	US-08-443-977-81	US-08-443-977-77	US-08-443-976-81
	Sequence 39, Appl	Sequence 1, Appli	Sequence 1, Appli	Sequence 44, Appl	Sequence 73, Appl	Sequence 65, Appl	Sequence 73, Appl	Sequence 65, Appl	Sequence 73, Appl	Sequence 65, Appl	Sequence 5, Appli	Sequence 34, Appl	•	Sequence 34, Appl	Sequence 33, Appl	Sequence 81, Appl	Sequence 77, Appl	Sequence 81, Appl

US-09-369-494-1 VS-09-369-494-1 VS-09-

Sequence 1, Application US/09569670

Patent No. 6689582

GENERAL INFORMATION:
APPLICANT: Davies, Christopher
APPLICANT: Chen, Davies, Steve
APPLICANT: Roczniak, Steve
TITLE OF INVENTION: Protein Having Proteinase Inhibitor Activity
FILE REFERENCE: MSB-7260
CURRENT APPLICATION NUMBER: US/09/569,670
CURRENT APPLICATION DAMBER: US/09/569,670
CURRENT FILING DATE: 2000-05-12
NUMBER OF SEQ ID NOS: 17
SOPTWARE: Patentin Ver. 2.0
SEQ ID NO 1
LENGTH: 58
TYPE: PRT

ORGANISM: Artificial Sequence

RESULT 2 US-09-569-670-1 밁

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1 YPURCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASBQBCMSSCQGS 58

YPVRCLLPSAHGSCADWAARWYFVASVGQCNRPWYGGCHGNANNFASEQECMSSCQGS 58

site Circuit

Matches

48;

Conservative

<u>,</u>

Mismatches

Indels

0

Gaps

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; OTHER INFORMATION: Description of Artificial Sequence: fragment from ; OTHER INFORMATION: computer database US-09-569-670-15
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Best Local S
Matches 48
                                                                                                                                                                                                                              SOFTWARE: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/369,494
CURRENT FILING DATE: 1999-08-05
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 15
LENGTH: 189
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Davies, Christopher
APPLICANT: Chen, Dadong
APPLICANT: Rozariak, Steve
TITLE OF INVENTION: Protein Having Proteinase Inhibitor Activity
FILE REFERENCE: MSB-7260
      Matches
                                            Query Match
                                                                                                                                                                                                                                             APPLICANT: Davies, Christopher
APPLICANT: Chen, Dadong
APPLICANT: Chen, Dadong
APPLICANT: Roczniak, Steve
TITLE OF INVENTION: Protein Having Proteinase Inhibitor Activity
FILE REFERENCE: MSB-7260
CURRENT APPLICATION NUMBER: US/09/569,670
CURRENT APPLICATION NUMBER: 2000-05-12
NUMBER OF SEQ ID NOS: 17
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15, Application US/09569670 Patent No. 6689582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15, Application US/09369494 Patent No. 6180607
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                                                                                                                                                                                        LENGTH: 189
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE: OTHER INFORMATION: Description of Artificial Sequence:fragment from OTHER INFORMATION: computer database
                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence:fragment from OTHER INFORMATION: computer database
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    Local Similarity
les 48; Conserv
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    Conservative
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82.8%; Pred. No. 5.9e-25;
vative 0; Mismatches 10
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80.2%; Score 264; DB 4;
82.8%; Pred. No. 5.9e-25;
tive 0; Mismatches 10
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Pred. No. 1.7e-25;
0; Mismatches 10
                                        Length 189
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US-09-369-494-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: MSB-7260
CURRENT APPLICATION NUMBER: US/09/369,494
CURRENT FILING DATE: 1999-08-05
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PACENTIN VET. 2.0
SEQ ID NO 9
LENGTH: 51
                                                                     Query Match
Best Local Similarity
Matches 42; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9, Application US/09569670 Patent No. 6689582 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                     SOFTWARE:
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/09369494 Patent No. 6180607
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                      FILE REFERENCE: MSB-7260
CURRENT APPLICATION NUMBER: US/09/569,670
CURRENT FILING DATE: 2000-05-12
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Chen, Dadong
APPLICANT: Roczniak, Steve
TITLE OF INVENTION: Protein Having Proteinase Inhibitor Activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Davies, Christopher
APPLICANT: Chen, Dadong
APPLICANT: Roczniak, Steve
TITLE OF INVENTION: Protein Having Proteinase Inhibitor Activity
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                              LENGTH: 51
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
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                                                                                                                                                                                                                                                                                           PatentIn Ver. 2.0
CLLPSAHGSCADWAARWYFVASVGQCNRFWYGGC-GNANNFASEQECMSSCQ
                        CLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQ 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLLPSAHGSCADWAARWYFVASVGQCNRFWYGGC-GNANNFASEQECMSSCQ 51
                                                                                                                                                                                                                                                                                                                                                                                                                                  Davies,
                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                           Description of Artificial Sequence: fragment from computer database
                                                                                                                                                                                                                                                                                                                                                                                                                                  Christopher
                                                                                      67.0%;
80.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       67.0%;
                                                                       0,
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                                                                   Score 220.5; DB 4;
Pred. No. 3.4e-20;
0; Mismatches 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 220.5;
Pred. No. 3.4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                       Indels
                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 51;
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 51
                                                                     Gaps
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RESULT 7 US-08-084-718-75

Sequence 75, Application US/08084718 Patent No. 5591603

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PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 4501/87
FILING DATE: 28 AUG-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 2254/88
FILING DATE: 26 APPR-1988
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DK88/00138
PRIOR APPLICATION NUMBER: PCT/DK88/00138
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DK89/00096
PRIOR APPLICATION NUMBER: PCT/DK89/00096
                                                                                                                                                                                                                                                                              PILING DATE: 12-JUN-1991

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DK91/00299

FILING DATE: 01-OCT-1991

PRIOR APPLICATION NUMBER: US 07/827,687

FILING DATE: 29-JAN-1992

PRIOR APPLICATION NUMBER: US 08/024,925

APPLICATION NUMBER: US 08/024,925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                     TELEFAX: 212-867-0298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Rele
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 19930623
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: No. 55916
APPLICANT: Petersen,
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 2361/90
FILING DATE: 01-OCT-1990
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                                      SEQUENCE CHARACTERISTICS
LENGTH: 109 amino aci
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 19-NOV-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 1118/91
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                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Agris, Cheryl H.
REGISTRATION NUMBER: 34,086
REFERENCE/DOCKET NUMBER: 311
FELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: US 07/598,337
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 07/466,408 FILING DATE: 21-JUN-1990
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FOPOLOGY:
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                        amino acid
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INVENTION: Process for the Preparation of Aprotinin
INVENTION: and Aprotinin Analogs
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Diness, Viggo
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                                                          FILING DATE: 19-NOV-PRIOR APPLICATION DATA:
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PRIOR APPLICATION NIMBER: US 07/466,408
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 2254/88
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           PRIOR APPLICATION DATA:
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                                                                                                                          APPLICATION NUMBER: DK 2361/90 FILING DATE: 01-OCT-1990
                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 26-APP
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                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 0 FILING DATE: 23-JUN-1993
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STATE: New Yor
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                                            APPLICATION NUMBER:
                                                                                           APPLICATION NUMBER:
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No. 5618915ris, Kjeld
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No. 5618915rskov-Lauritsen,
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63.0%;
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                                                                                                                                                                                                                                        PCT/DK89/00096
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                                                                                           US 07/598,337
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PCT/DK91/00299
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                                            DK 1118/91
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Pred. No. 2.6e-18;
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US 07/827,687

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                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/443,977
FILING DATE: 18-MAY-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 75:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US 07/1
PILING DATE: 29-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/1
FILING DATE: 26-FEB-1993
ATTORNEY/AGENT INFORMATION:
NAME: AGIST, Cheryl H.
REGISTRATION NUMBER: 34,08
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TOPOLOGY: 1:-
                                APPLICATION NUMBER: DK 4501/87
FILING DATE: 28-AUG-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 2254/88
FILING DATE: 26-APR-1988
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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ADDRESSEE: No. 56210
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TELEPHONE: 212-867-0298
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: United States of America
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No. 5621074rs, Niels D.
Christensen, Niels D.
Bregengaard, Claus
No. 5621074ris, Fanny
Petersen, Lars C.
Petersen, Lars C.
Process for the Preparation of Aprotinin
Analogs
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No. 5621074ris, Kjeld
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ilarity 63.0%;
Conservative
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PCT/DK88/00138
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; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-443-977-75
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Patent No.
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Best Local Similarity 63.0%;
Matches 34; Conservative
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INFORMATION FOR SEQ ID NO:
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APPLICANT:
APPLICANT:
                                                                                                                                                          APPLICANT: Christensen, Niels D.
APPLICANT: Bregengaard, Claus
APPLICANT: No. 5591003ris, Fanny
APPLICANT: Petersen, Lars C.
TITLE OF INVENTION: Process for the Preparation of Aprotinin
TITLE OF INVENTION: and Aprotinin Analogs
NUMBER OF SEQUENCES: 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
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NAME: AGENT INFORMATION:
REGISTRATION NUMBER: 34,0
REFERENCE/POOPURE: 34,0
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APPLICATION NUMBER: PCT/L
FILING DATE:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                           CORRESPONDENCE ADDRESS
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PRIOR APPLICATION NIMBER: DK 1118/91
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PRIOR APPLICATION DATA:
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                                  COUNTRY: United States of America ZIP: 10174-6201
                                                                                          CITY: New York
                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 34,086
REFERENCE/DOCKET NUMBER: 31
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FILING DATE: 25-APR-1989
                                                                                                                             ADDRESSEE:
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5591603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  56 CLEPPSTGPCKARIIRYFYNAKAGLCQTFVYGGCRGNGNNFKSAEDCMETCGGA 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
                                                                           New York
                                                                                                                                                                                                                                                                                                                                                                                               Application US/08084718
                                                                                                            405 Lexington Avenue,
                                                                                                                                                                                                                                                                                       Diness, Viggo
No. 5591603rskov-Lauritsen, Leif
                                                                                                                                                                                                                                                                                                                     Bjorn, Soeren E.
No. 5591603ris, Kjeld
                                                                                                            No. 55916030 No. 5591603disk of No. Stexington Avenue, 62nd Floor
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Pred. No. 2.6e-18;
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                                                                                                                             5591603th America,
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; MOLECULE TYPE: protein
US-08-084-718-59
                                                       Sequence 59, Applicati
Patent No. 5618915
GENERAL INFORMATION:
APPLICANT: Bjorn,
                                                                                                                                                                                                                                                                            Query Match 61.7%;
Best Local Similarity 59.6%;
Matches 34; Conservative
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                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
LENGTH: 111 amino aci
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER:
FILING DATE: 01-OCT-
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICATION NUMBER: US 07/598,337
FILING DATE: 19-NOV-1990
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APPLICATION NUMBER: 1
FILING DATE: 25-APR-1
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APPLICATION NUMBER: DK 4501/87
FILING DATE: 28-AUG-1987
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
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APPLICATION NUMBER:
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APPLICATION NUMBER:
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NTTORNEY/AGENT INFORMATION:
NAME: Agris, Cheryl H.
REGISTRATION NUMBER: 34,0
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                                                                                                                                                                                                       55 PDFCLEPPSTGPCKARIIRYFYDATAGLCETFVYGGCRANRNNFKSAEDCMETCGGA 111
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                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
                                                                                                             Application US/08443976
Bjorn, Soeren E.
No. 56.8915r1s, Kjeld
Diness, Viggo
No. 5618915rskov-Lauritsen, Leif
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21-JUN-1990
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                                                                                                                                                                                                                                                                          Score 203; DB 1; Length 111; Pred. No. 1.1e-17; 9; Mismatches 14; Indels
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; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-443-976-59
                                                                                                   TELEFAX: 212-867-0298
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PRIOR APPLICATION DATA:
PRIOR TAMETON NUMBER: US 08/024,925
                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/DK91/00299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 21-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 2361/90
                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                         FILING DATE: 01-OCT-1991 PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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APPLICATION NUMBER: PCT/DK89/00096
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 2254/88
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TITLE OF INVENTION: Process for the Preparation of Aprotinin
TITLE OF INVENTION: and Aprotinin Analogs
NUMBER OF SEQUENCES: 83
                                                                                                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0
                                                TYPE: amino acid
                                                                                                                                                                                               NAME: Agris, Cheryl H. REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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CITY: New York
                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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                                                                       111 amino acids
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                                                                                                                                            212-867-0123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DK 4501/87
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                                                                                                          59:
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Query Match Best Local Similarity

61.7**%**; 59.6**%**;

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RESULT 12
US-08-443-977-59
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Sequence 59, Appro-
No. 562107/
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                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 530
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/084,718
FILING DATE: 23-UN-1993
APPLICATION NUMBER: DK 4501/87
FILING DATE: 28-AUG-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 2254/88
FILING DATE: 26-APR-1988
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DK88/00138
FILING DATE: 28-AUG-1988
PRIOR APPLICATION NUMBER: PCT/DK89/00096
APPLICATION NUMBER: PCT/DK89/00096
APPLICATION NUMBER: PCT/DK89/00096
APPLICATION NUMBER: PCT/DK89/00096
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                                                         APPLICATION NUMBER: DK 1:
FILING DATE: 12-JUN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/I
FILING DATE: 01-CCT-1991
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ZIP: 10174-6201

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/443,977
FILING DATE: 18-MAY-1995
                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/598,337
FILING DATE: 19-NOV-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 07/466, FILING DATE: 21-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 2361/90
FILING DATE: 01-OCT-1990
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APPLICANT: Petersen, Lars C.
TITLE OF INVENTION: Process for the Preparation of Aprotinin
TITLE OF INVENTION: and Aprotinin Analogs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 PVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08443977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: No. 5621074o No. 5621074disk of No. 5621074th America,
405 Lexington Avenue, 62nd Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              No. 5621074rskov-Lauritsen, Leif
Christensen, Niels D.
Bregengaard, Claus
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No. 5621074ris, Kjeld
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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MBER: US 07/827,687
29-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCT/DK88/00138
                                                                                   PCT/DK91/00299
                                                                                                                                                        DK 1118/91
                                                                                                                                                                                                                                                                                                                                                             US 07/466,408
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-084-718-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 71,
Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 61.7%;
Best Local Similarity 59.6%;
Matches 34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 212-867-0298
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: NO. 55910
APPLICANT: Petersen.
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                  APPLICATION NUMBER: UP11 DATE: 19930623 CLASSIFICATION: 514 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF CONTRESPONDENCE ADDRESS:

ADDRESSEE: No. 55916030 No. 5591603disk of No. ADDRESSEE: No. 55916030 No. 62nd Floor
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APPLICANT:
APPLICANT:
                                                 PILING DATE: 28-AUG-1988
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/
                                                                                                                                                                             APPLICATION NUMBER: DK 4501/87 FILING DATE: 28-AUG-1987 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 111 amino acids
                                                                                                                          FILING DATE: 26-APR-1988 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
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                                    FILING DATE:
                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: New York
STATE: New York
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REGISTRATION NUMBER:
                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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VENTION: Process for the Preparation of Aprotinin
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No. 559160378kov-Lauritsen, Leif
Christensen, Niels D.
Bregengaard, Claus
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                                  25-APR-1989
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                                                                                                                                                                DK 2254/88
                                                     PCT/DK89/00096
                                                                                                        PCT/DK88/00138
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US 07/466,408
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Pred. No. 1.1e-17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fanny
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21-JUN-1990

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APPLICATION NUMBER: US 07/827

FILING DATE: 29-UAN-1992

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/024

FILING DATE: 26-FEB-1993

ATTORNEY/AGENT INFORMATION:

NAME: AGRIS, Cheryl H.

REGISTRATION NUMBER: 31,086

REFERENCE/DOCKET NUMBER: 3143

TELECOMMUNICATION INFORMATION:

TELEPHONE: 212-867-0123

TELEPHONE: 212-867-0123
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 56189
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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APPLICATION NUMBER:
FILING DATE: 01-OCT-
                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                    SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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APPLICATION NUMBER:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                            STATE: New York
    APPLICATION NUMBER:
                                                                                                                                                                                              CITY: New York
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                                                                                                                                                        COUNTRY:
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                                                                                                                                                        United States of America
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                                                                                                                                                                                                                                                                                         No. 5618915ris, Fanny
Petersen, Lars C.
MENTION: Process for the Preparation of Aprotinin
ENTION: and Aprotinin Analogs
                                                                                                                                                                                                                                                                                                                                                                     Christensen, Niels D. Bregengaard, Claus
                                                                                                                                                                                                                                                                                                                                                                                                           Diness, Viggo
No. 5618915rskov-Lauritsen, Leif
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No. 5618915ris, Kjeld
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                                                                                                                                                                                                                                No. 56189150 No. 5618915disk of No. 5618915th America,
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01-OCT-1991
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US/08/443,976
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                                                                                                                                                                                                                  62nd Floor
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Sequence 71, Application Patent No. 5621074
GENERAL INFORMATION:
APPLICANT: Bjorn,

Application US/08443977

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APPLICANT: APPLICANT:

APPLICANT:

Christensen, Niels D. Bregengaard, Claus No. 5621074ris, Fanny

Bjorn, Soeren E. No. 5621074ris, Kjeld Diness, Viggo No. 5621074rskov-Lauritsen,

Soeren E.

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Query Match
Best Local Similarity
Matches 33; Conserve
                                                                                                                                                                                              ; MOLECULE TYPE: protein US-08-443-976-71
RESULT 15
US-08-443-977-71
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: DK 11
FILING DATE: 12-UTW-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/I
FILING DATE: 01-CCT-1991
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APPLICATION NUMBER: DK 1118/91
FILING DATE: 12-UUN-1991
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                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Agris, Cheryl H.
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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FILING DATE: 26-APR-
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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CLASSIFICATION: 530
PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: DK 4501/87
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                                                                  56 CLEPPSTGPCKARIIRYFYDATAGLCETFVYGGCRANRNNFKSAEDCMETCGGA 109
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                                                                                                                                                                                                                                amino acid
GY: linear
                                                                                   CLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                             109 amino acids
                                                                                                                                  Conservative
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26-APR-1988
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                                                                                                                                              61.4%;
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                                                                                                                                                                                                                                                                                                                                                                               34,086
                                                                                                                                                                                                                                                                                                  71:
                                                                                                                                                                                                                                                                                                                                                                 3143.224-US
                                                                                                                                 9
                                                                                                                                 Score 202; DB 1;
Pred. No. 1.4e-17;
9; Mismatches 12
                                                                                                                                                              Length 109;
                                                                                                                                 Indels
                                                                                                                                 ٥,
                                                                                                                                 Gaps
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; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-443-977-71
Query Match 61.4%; Score 202; DB 1; Length 109; Best Local Similarity 61.1%; Pred. No. 1.4e-17; Matches 33; Conservative 9; Mismatches 12; Indels
                                                                                                                                                                                                                                                                                                                            APPLICATION UNMBER: US 07/466,408
FILING DATE: 21-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION UNMBER: DK 2361/90
PILING DATE: 01-OCT-1990
PRIOR APPLICATION UNMBER: US 07/598,337
FILING DATE: 19-NOV-1990
PRIOR APPLICATION UNMBER: DK 1118/91
FILING DATE: 12-JUN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DK91/00299
FILING DATE: 01-OCT-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DK91/00299
FILING DATE: 01-OCT-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/827,687
FILING DATE: 29-JAN-1992
PRIOR APPLICATION NUMBER: US 08/024,925
FILING DATE: 26-FEB-1993
ATTORNEY/AGENT INFORMATION:
NAME: AGTIS, Cheryl H.
REGISTRATION NUMBER: 34,086
REFERENCE/DOCKET NUMBER: 3143.224-US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 08/084,718

RIPLICATION NUMBER: US 08/084,718

RILING DATE: 23-UN-1993

APPLICATION UNMBER: DK 4501/87

FILING DATE: 28-AUG-1987

PRIOR APPLICATION DATA:

APPLICATION NUMBER: DK 2254/88

FILING DATE: 26-APR-1988

FILING DATE: 26-APR-1988

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/DK88/00138

FILING DATE: 28-AUG-1988

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/DK89/00096

FILING DATE: 25-APR-1989

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/466,408

APPLICATION NUMBER: US 07/466,408
                                                                                                                                                                                                                                                    TELEFAX: 212-867-0298
INFORMATION FOR SEQ ID NO: 71:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNMBER: US/08/443,977
FILING DATE: 18-MAY-1995
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: New York
COUNTRY: United States of America
ZIP: 10174-6201
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 8:
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TITLE OF INVENTION: Process for the Preparation of Aprotinin
TITLE OF INVENTION: and Aprotinin Analogs
                                                                                                                                                                                                LENGTH:
TYPE: a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 405 Le:
CITY: New York
STATE: New Yor!
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                                                                                                                                                                                                                    109 amino acids
                                                                                                                                                                                                                                                                                                          212-867-0298
                                                                                                                                                                                                                                                                                                                                                                                              3143.224-US
  0,
Gaps
     0,
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Search completed: August 25, 2005, 04:56:24 Job time : 33 secs

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Title:
Perfect score:
Sequence:
                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OM protein - protein search, using sw model
                                                                                                                                                             Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                                                                                    Searched:
                                                                                                                                                                                                                                                                                                                         Scoring table:
                                                 Database :
                                                                                                                                                                                                                                                              1612378 seqs, 512079187 residues
                                                                                                                                                                                                                                                                                                     BLOSUM62
Gapop 10.0 , Gapext 0.5
UniProt_03:*
1: uniprot_sprot:*
2: uniprot_trembl:*
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329
1 YPVRCLLPPATGPCKARIIR.....RGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                            August 25, 2005, 04:45:06; Search time 176 Seconds (without alignments)
168.753 Million cell updates/sec
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Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                               1612378
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

30	26 28 28	222 223 254	16 17 19	13 15	110 121 121	4664D06	Result No.
165	166 166 166	1688 1688 1688 1688	172 168 168	184 177 174	188 186 186 185	264 264 264 192	Score
	, , , , , , , , , , , , , , , , , , ,	51.1 51.1 51.1	51.1 51.1			5 7 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	Query Match 1
252 252	576 83	738 751 763 763	80 111 523 738	100 82 122	984 984 327	531 659 660 1235 1280 100	Length D
		004004 004004				QQQQQEQ QQQQQEQ	DB ID
Q90WAO SPT2_HUMAN	Q6ZN14 Q8TEU8 Q6UXZ9	Q60709 Q60709 APP2 HUMAN Q71UIO Q61482 APP2 RAT	Q8T3S7 Q7M4L3 Q14594 Q90W28	BPT2_BOVIN Q7ZZT3 UPTI_PIG	Q7M312 Q9GQN1 Q9GQN2 Q6IND9 IBPS BOVIN	Q659F2 Q6ZMM1 Q7Z681 Q7Z681 Q9E9EX2 Q9EPX2 BPT1 BOVIN Q7M311	•
Q90wa0 Q43291			Q8t3s7 Q7m4l3 Q14594 Q90w28		0.00.00	Q659f2 Q6znml Q7z681 Q9z681 Q9epx2 Q9epx2 P00974 Q7m31	Description
pseudonaja homo sapien	nomo sapien homo sapien homo sapien	brachydanio mus musculu homo sapien homo sapien mus musculu rattus norv	araneus ven homo sapien homo sapien brachydanio	Ω	0	homo sapien homo sapien homo sapien homo sapien homo sapien mus musculu bos taurus bos indicus	

164 49.8 61 1 VIDAN 164 49.8 90 2 Q6XPV8 163 49.5 57 2 Q7M4I 163 49.5 62 1 IVBT OXYSC 163 49.5 110 1 IBP CARCR 163 49.5 110 1 IBP CARCR 163 49.5 193 2 Q6P796 163 49.5 193 2 Q7TQN3 162 49.2 58 1 AXP1 ANTAF 162 49.2 58 1 AXP1 ANTAF 162 49.2 58 2 Q9TWF9 162 49.2 58 2 Q9TWF9 162 49.2 58 3 Q9TWF9 163 48.9 133 2 Q8BZ45 161 48.9 253 2 Q16701 161 48.9 751 2 Q9DGJ7	49.8 61 1 1 49.8 57 2 2 49.5 62 1 1 1 49.5 110 2 1 49.5 571 2 2 49.2 58 2 2 2 49.2 58 2 2 2 49.2 58 2 2 2 49.2 58 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	45	44	43	42	41	40	39	38	37	36	3 5	34	33	26
61 1 1 90 2 0 50 2 0 62 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	61 I VERA POOSIO 61 I VERA POOSIO 61 I VERA POOSIO 90 2 QAXPYB 91 2 QAMAII 162 I IVBT QAXXSC 110 1 IBP CARCR 100 1 IBP CARCR 100 2 QAF796 193 2 QAF796 193 2 QAF797 198 2 QAYRAF 19	161	161	161	162	162	162	163	163	163	163	163	164	164	FOO
	1 IVB1 VIPAA P00991 2 Q6XFY8 Q6XFY8 2 Q7M411 1 IVBT QXYSC Q712e4 1 IBP CARCR P00993 2 Q6F796 Q7FQm3 2 Q7TQN3 Q7FQm3 2 Q9TWF9 Q9FWF9 2 Q9M98 Q9W98 2 Q9M93 Q9BT245 2 Q16T701 Q9dgj7 2 Q9DGJ7 Q9dgj7	48.9	48.9	48.9	49.2	49.2	49.2	49.5	49.5	49.5	49.5	49.5	49.8	49.8	00.1
00000200440044	VIDAA P00991 VB1 VIDAA P00991 VB1 VIDAA P00991 Osapy8 Oracin Osapy8 Oracin O	751	253	133	83	58	58	571	193	110	62	57	90	61	404
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		Q9DGJ7	016701	Q8HZ45	Q90W98	Q9TWF9	AXP1_ANTAF	Q7TQN3	Q6P796	IBP_CARCR	IVBT_OXYSC	Q7M4I1	Q6XPY8	IVB1_VIPAA	Corpo

ALIGNMENTS

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A Wambutt R., Heubner D., Mewes H.W., Weil B., Amid C., Osang A Fobo G., Han M., Wiemann S.;
L Submitted (SEP-2004) to the EMBL/GenBank/DDBJ databases.
C -!- SIMILARITY: Contains 1 BPTI/Kunitz inhibitor domain.
R EMBL; BX537757; CAD97826.1; -.
R HSSP; Q16019; 1AAP.
NR HSSP; Q16019; 1AAP.
GO; GO:0004867; F:serino-type endopeptidase inhibitor activ InterPro; IPR007110; Ig-1ike.
DR InterPro; IPR003598; Ig_C2.
DR InterPro; IPR003598; Frot_Inh_Kunz-m.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 48
                                                                  Pfam; PF00047; 1g; 1.

Pfam; PF00014; Kunitz BPTI; 1.

ProDom; PF00022; Prot Inh Kunz-m; 1.

SMART; SM00408; IGC2; 3.

SMART; SM00131; KU; 1.

PROSITE; PS00280; BPTI KUNITZ 1; 1.

PROSITE; PS5035; IG LIKB; 3.

PROSITE; PS50900; PLAC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pfam; PF00047; 14; 1.

Pfam; PF00014; Kunitz BPTI; 1.

Pfam; PF00018; Kunitz BPTI; 1.

PRINTS; PR00759; BASICPTASE.

ProDom; PD000222; Prot Inh_Kunz-m; 1.

SMART; SM00409; IG; 3.

SMART; SM00406; IGc; 3.

SMART; SM00406; IGc; 3.

SMART; SM00406; IGv; 3.

SMART; SM00131; KU; 1.

PROSITE; PS00230; BPTI KUNITZ_1; 1.

PROSITE; PS00230; BPTI KUNITZ_2; 1.

PROSITE; PS50935; IG_LIKE; 3.

PROSITE; PS50930; PLAC; 1.

NON TER

SEQUENCE 659 AA; 70274 MW; 50FFAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HSSP; Q16019; 1A
GO; GO:0004867;
InterPro; IPR003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q7Z681 PRELIMINARY; PRT; 660 AA.
Q7Z681;
Q1-OCT-2003 (TrEMBLrel. 25, Created)
O1-OCT-2003 (TrEMBLrel. 25, Last sequence update)
O1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Hypothetical protein DKFZp779D0163 (Fragment).
Name=DKFZp779D0163;
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InterPro;
InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human).
                      Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9606;
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); IPR003598; Ig-c2.

); IPR003596; Ig-c2.

); IPR010909; PLAC.

); IPR0102223; Prot_Inh_Kunz-m.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 188
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660
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ΑA,
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70374 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                80.2%;
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Pred. No. 3.1e-23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
A6D5B1C50CBB8815 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              50FFA993BB44C092 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                             inhibitor activity; IEA.
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CKESULT 5
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Best Local S
Matches 48
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Best Local Similarity
Matches 48; Conserv
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095428;
095428;
01-MAY-1999 (TEMBLrel. 1
01-MAY-1999 (TEMBLrel. 1
01-MAR-2004 (TEMBLrel. 2
                                                                   Q9EPX2;
Q9EPX2;
~01-MAR-2001
01-MAR-2001
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF05986; ADAM spacer1; 1.
Pfam; PF00047; 19; 1.
Pfam; PF00014; KUnitz BPTI; 1.
Pfam; PF00090; TSP 1; 5.
PFNINTS; PR00759; BASICPTASE.
PRODOM; PD000222; Prot Inh_Kunz-m; 1
SMART; SM00408; IGC2; 3.
SMART; SM00131; KU; 1.
SMART; SM00209; TSP1; 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE; PS50
PROSITE; PS50
Hypothetical
SEQUENCE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rowen L., Madan A., Qin S., Abbasi N., Dors M., Ratcliffe Madan A., Dickhoff R., Shaffer T., James R., Lasky S., Hoc Submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Contains 1 BPTI/Kunitz inhibitor domain.
EMBL; AF109907; AAC97963.1; -.
HSSP; P12111; 1KTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hypothetical protein.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
NCBI_TaxID=9606;
                      Name=Papln;
                                          Papilin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PS00280; BPTI_KUNITZ_1;
PROSITE; PS50279; BPTI_KUNITZ_2;
PROSITE; PS50835; IG_LIKE; 3.
PROSITE; PS50900; PLAC; 1.
PROSITE; PS50902; TSP1; 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR010294; ADAM spacer1.
InterPro; IPR007110; Ig-1ke.
InterPro; IPR003598; Ig_c2.
InterPro; IPR010909; PLAC.
InterPro; IPR0002223; Prot Inh_Kunz-m.
InterPro; IPR000884; TSP1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GO; GO:0004867; F:serine-type endopeptidase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genew; HGNC:19262; PAPLN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
  musculus
                                                                                                                                                                                                                                                                                        883
                                                                                                                                                                                                                                                                                                                                                                                  48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS
                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                 YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                      YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           al protein.
1235 AA; 1
                                                                                                                                                                                                                                                                                                                                                                               nilarity (80.2%;)
Conservative 0
                                                                     (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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                                                                                                                                                                    PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
(Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                133476 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               80.2%;
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26,
                                                                        Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                               Score 264; DB 2; Pred. No. 5.6e-23; 0; Mismatches 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Created)
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Pred. No. 3.1e-23;
0; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Craniata; Vertebrata;
Catarrhini; Hominidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A0B44CCE4F38E350 CRC64;
                                                                                                                                                                 Å
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hominidae;
                                                                     update)
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                                                                                                                                                                                                                                                                                                                                                                                                                               Length 1235;
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Best Local S
Matches 43
                            protease
J. Mol. B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BPT1_BOV
P00974;
                                                     SEQUENCE FROM N.A.

MEDLINE-87283904; PubMed=2441071;

Creighton T.E., Charles I.G.;

"Sequences of the genes and polypeptide
protease inhibitors.";
                                                                                                                                                                                                                                                                                                                                                                                                             21-JUL-1986 (Rel. 01, Created)
01-MAR-1989 (Rel. 10, Last sequence upd
25-OCT-2004 (Rel. 45, Last annotation u
Pancreatic trypsin inhibitor precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROSITE; PS00280; BPTI KUNITZ 1; 1.
PROSITE; PS50279; BPTI KUNITZ 2; 1.
PROSITE; PS50835; IG LIKE; 3.
PROSITE; PS50879; ODR DC 2; UNKNOWN 1.
PROSITE; PS50900; PLAC; 1.
PROSITE; PS50092; TSP1; 5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam;
Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sieron A.L., Prockop D.J., Fessler J.H.;
"Papilin in development; a pericellular protein
ADAMTS metalloproteinasses.";
Development 127:5475-5485(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BOVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ProDom; PD000222; Prot
SMART; SM00408; IGc2;
                                                                                                                                                                                                                                                                         Bovinae; Bos
                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                              Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRINTS; PR00759; BASICPTASE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ဗိ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kramerova I.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                    NCBI_TaxID=9913;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- SIMILARITY: Contains 1 BPTI/Kunitz
EMBL; AF314171; AAG41980.1; -.
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[1]
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Mammalia; 1
2
                                                                                                                                                                                                                                                                                                                                                                                             (BPI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro;
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DO; GC:0004867; F:serine-type endopeptidase in:
nterPro; IPR010294; ADAM_spacer1.
nterPro; IPR000183; Decarbxy1se2.
nterPro; IPR007110; Ig-like.
nterPro; IPR003598; Ig_C2.
nterPro; IPR0103909; PLAC.
nterPro; IPR010909; PLAC.
nterPro; IPR0102223; Prot Inh_Kunz-m.
nterPro; IPR000884; TSP1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P12111; 1KTH
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SM00131; KU; 1.
SM00209; TSP1; 5
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                                  Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Metazoa;
Eutheria;
                                                                                                                                                                                                                                                                                                                                                           (Bovine)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1280
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                                  194:11-22(1987)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kunitz_
TSP 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
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                                                                                                                                                                                                                                                                                               Chordata; Craniata; Vertebrata; Euteleostomi; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chordata;
Rodentia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    138823 MW;
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4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 246;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
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J.M., Ackley B.D.,
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MEDLINE-91332906; Pubmeu---- WOO Housset D., Kim K.-S., Fuchs J., Woo Housset D., Kim K.-S., Fuchs J., Woo
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MEDLINE=90211226; PubMed=2322242;
Lewis R.V., Ray P., Coguill R., K
"Presence of pancreatic trypsin in
chromaffin cells.";
                                                                                                                                                           Deisenhofer J., Steigemann W.;
"Crystallographic refinement of the structure
trypsin inhibitor at 1.5-A resolution.";
Acta Crystallogr. B 31:238-250(1975).
                                                                                                                                                                                                                                                                                                                                     SEQUENCE OF 36-93.
Dlouha V., Pospisilova
"Sequence of residues 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Isolation of a genomic clone for bovine pancreatic by using a unique-sequence synthetic DNA probe."; proc. Natl. Acad. Sci. U.S.A. 80:6838-6842(1983).
                                                                             Huber R., Kukla D., Ruhlmann A., Epp O., Formanel "The basic trypsin inhibitor of bovine pancreas. and conformation of the polypeptide chain."; Naturwissenschaften 57:389-392(1970).
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                                                                                                                                                                                                                                Biochem. Biophys. Res. Commun. 167:543-547(1990)
                                                                                                                                                                                                                                                                                                    SEQUENCE OF 36-81.
                                                                                                                                                                                                                                                                                                                                                                                              "Covalent structure of a polypeptide inhibitor Northrop inhibitor).";
                                                                                                                                                                                                                                                                                                                                                                                                                     Chauvet J., Acher R.;
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MEDLINE=68012003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     J. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anderer F.A., Hornle S.; "The disulfide linkages
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kassell B., Laskows
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inhibitor.";
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                                                      X-RAY CRYSTALLOGRAPHY (1.8 ANGSTROMS)
                                                                                                                                      X-RAY CRYSTALLOGRAPHY (2.5 ANGSTROMS).
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220:757-770(1991)
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EMBL; M20934; AAD13685.1; -.
EMBL; M20930; AAD13685.1; JOINED.
EMBL; M20932; AAD13685.1; JOINED.
EMBL; X03365; CAA27062.1; ALT_SEQ.
EMBL; X03365; CAA27063.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Berndt K.D., Guntert P., Orbons L.P.M., Wuethrich K., "Determination of a high-quality nuclear magnetic restructure of the bovine pancreatic trypsin inhibitor with three crystal structures.";
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STRUCTURE BY NMR.
STRUCTURE BY NMR.
MEDLINE=93021116; PubMed=1383552;
MEDLINE=93021116; PubMed=1383552;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      surgery.
SIMILARITY: Contains 1 BPTI/Kunitz inhibitor domain DATABASE: NAME=Trasylol; NOTE=Clinical information WWW-"http://www.trasylol.com/".
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mol. Biol. 227:757-775(1992).
FUNCTION: Inhibits trypsin, kallikrein,
SUBCELLULAR LOCATION: Secreted.
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10A6;
1PIT;
1QLQ;
1TPA;
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Siekmann J., Wenzel H.R., Schroeder W., Tschesche H.;

Siekmann J., Wenzel H.R., Schroeder W., Tschesche H.;

Siekmann J., Wenzel H.R., Schroeder W., Tschesche H.;

Thomologues from bovine lungs ";

Biol. Chem. Hoppe-Seyler 369:157-163(1988).

-!- SIMILARITY: Contains 1 BPTI/Kunitz inhibitor domain.

PIR, S00371, S00371.

R PIR, S00371, 1BPI.

R HSSP; P00974; 1BPI.

R GO; GO:0004867; F:serine-type endopeptidase inhibitor active gram; pro10014; Kunitz BPTI; 1.

Pfam; pF00014; Kunitz BPTI; 1.

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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
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PRINTS; PR00759; BASTCPTASE.

ProDom; PD000222; Prot_Inh_Kunz-m;

SMART; SM00131; KU; 1.

PROSITE; PS00280; BPTI_KUNITZ 1; 1

PROSITE; PS50279; BPTI_KUNITZ_2; 1
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Potential.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 192; DB Pred. No. 2.3e. 7; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           trypsin inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
.3e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ₿
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pharmaceutical;
                                                                                                   inhibitor activity;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 100;
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                                                                                                                                                                                                                 aprotinin
                                                                                                                                                                                                                                                                                                                                                          Euteleostomi;
cora; Bovidae;
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                                                                                                   IEA
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Q9GQN1
ID Q9GQ
AC Q9GQ
AC Q9GQ
AC Q9GQ
DT 01-M
DT 01-G
Pept
GN Name
OS Call
OC Nyna
OX NCBI
RN [1]
RP SEQU
RA Will
CCC --!-
DR EMBH
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O7M312
ID Q7
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RRP SB8
OC B8
OC B8
OC M8
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Best Local S
Matches 32
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                      Q9GQN1
Q9GQN1;
01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROSITE;
SEQUENCE
Williamson M., Hauser F., Grimmelikhuij
Submitted (JAN-2000) to the EMBL/GenBan
-!- SIMILARITY: Contains 1 BPTI/Kunitz
EMBL; AF221986; AAG44251.1; -.
HSSP; P12111; 1KNT.
                                                                                                                                                                                  Calliactis parasitica (Sea anemone).
Eukaryota; Metazoa; Cnidaria; Anthozo
Nynantheae; Hormathiidae; Calliactis
                                                                                                                                                                                                                                                                                                          01-MAR-2001
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRODOM; PD000222; Prot_Inh_Kunz-m; 1.
PROSITE; PS00280; BPTI_KUNITZ_1; 1.
PROSITE; PS50279; BPTI_KUNITZ_2; 1.
SEQUENCE 58 AA; 6532 MW; B858E86092EA1518 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PIR; S10063; S10063.

HSSP; P00974; IBPI.

GO; GO:0004867; P:serine-type endopeptidase inhibitor activity; IEA.

InterPro; IPR002223; Prot Inh Kunz-m.

Pfam; PF00014; Kunitz BPTT; 1.

PRINTS; PR00759; BASICPTASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     homologues from bovine lungs.";
Biol. Chem. Hoppe-Seyler 369:157-163(1988)
-!- SIMILARITY: Contains 1 BPTI/Kunitz inh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q7M312;
01-MAR-2004
                                                                                                                                                                                                                                                                                  Peptidylglycine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Siekmann J., Wenzel H.R., Schroeder W., Characterization and sequence determina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isoaprotinin G2.
Bos indicus x Bos taurus (hybrid cattle)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q7M312
                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                               NCBI_TaxID=6114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=88221840; PubMed=2453200;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=30522;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PS50279; BPTI_KUNITZ_2; 1.
59 AA; 6591 MW; B983C93192BB4418 CRC64;
                                                                                                                                                                                                                                                                           (TrEMBLrel. 16, Created)
(TrEMBLrel. 16, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
ine alpha-amidating monooxygenase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cetartiodactyla; Ruminantia; Pecora; Bovidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 57.8%;
56.1%;
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56.1%;
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                                                Grimmelikhuijzen C.J.P.;
he EMBL/GenBank/DDBJ databases.
1 BPTI/Kunitz inhibitor domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9;
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Pred. No. 2.3e-15
9; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. 4e-1
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 188;
                                                                                                                                                                                                            Anthozoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             determination
                                                                                                                                                                                                                                                                                                                                                                                                   984
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inhibitor domain
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                                                                                                                                                                                                            Zoantharia; Actiniaria;
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PROBLEM TO THE PROBLE
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UNITED AGAINST BETTI.
UNITED AGAINST
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Best Local
                                                                                      InterPro; IPR001258; NHL.
InterPro; IPR000720; Pamonoxygenase.
InterPro; IPR000720; Pamonoxygenase.
InterPro; IPR008977; PHM PNGase_F.
InterPro; IPR002223; Prot_Inh_Kunz-m.
Pfam; PF00182; Cu2_monoxygen; 1.
Pfam; PF03712; Cu2_monox C; 1.
Pfam; PF00014; Kunītz_BPTī; 1.
Pfam; PF00014; Kunītz_BPTī; 1.
Pfam; PF0001436; NHL; 5.
PRINTS; PR00759; PAMONOXGNASE.
PRINTS; PR00759; PAMONOXGNASE.
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01-MAR-2001
01-MAR-2001
01-OCT-2003
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InterPro;
                                    ProDom;
SMART; S
                                                                                                                                                                                                                                                                                                                                                                   GO; GO:0016020; C:membrane; IEA.
GO; GO:000520; F:copper ion binding; IEA.
GO; GO:0004504; F:poptidylglycine monooxygenase activity; IEA.
GO; GO:0004504; F:serine-type endopeptidase inhibitor activity;
GO; GO:0006518; P:septide metabolism; IEA.
InterPro; IPR011044; Amine DH B like.
InterPro; IPR000323; CU2 monooxygnse.
InterPro; IPR001258; NHL.
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Pfam;
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Pfam;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Williamson M., Hauser F., Grimmelikhuijzen C.J.
Submitted (JAN-2000) to the EMBL/GenBank/DDBJ d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=6114;
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Eukaryota; Metazoa; Cnidaria; Anthoz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Calliactis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptidylglycine alpha-amidating monooxygenase
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PROSITE; PS50279; BPTI_KUNITZ_2; 1.
PROSITE; PS00084; CUZ_MONOOXYGENASE_1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRINTS; PR00790; PAMONOXGNASE. SMART; SM00131; KU; 1.
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InterPro; IPR002223; Prot Inh Kunz-m.
Pfam; PPf0182; Cu2_monooxygen; 1.
Pfam; PF03712; Cu2_monoox_C; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro;
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GO:0005507; F:copper ion binding; IEA.
GO:0004504; F:peptidylglycine monooxygenase activity;
GO:0004667; F:serine-type endopeptidase inhibitor acti
GO:0004667; F:seride metabolism; IEA.
GO:0006518; P:peptide metabolism; IEA.
erpro; IPR011044; Amine_DH_B_like.
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                                 SM00131;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30;
                             PD000222; Prot_Inh_Kunz-m;
SM00131; KU; 1.
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IPR001258; NHL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hormathiidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (TrEMBLrel.
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BPTI_KUNITZ_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   110265 MW; 4C24E4BCEB3B480D CRC64;
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REDLINE=2238257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RICAL STRAUSDER R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altgebrul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altgebrul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,

A Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,

A Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,

A Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hsieh F.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Robak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Pahey J., Helton E., Ketteman M., Madan A., Godrigues S., Sanchez A.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Holkesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Bakesley R.W., Touchman J.W., Green E.D., Schnerch A., Schein J.E.,

Bakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Bakesley R.W.,
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Q6IND9;
05-JUL-2004
Klein S., Strausberg R.;
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Contains 3 BPTI/Kunitz inhibitor domains.
EMBL; BC072344; AAH72344.1; -.
HSSP; Q16019; 1AAP.
GO; GO:0005576; C:extracellular; IEA.
GO; GO:0030414; F:protease inhibitor activity; IEA.
GO; GO:0004867; F:serine-type endopeptidase inhibitor act
GO; GO:0007596; P:blood coagulation; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jones S.J., Marra M.A.;
Jones S.J., Marra M.A.;
Generation and initial analysis of more than 15,09
and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002)
                                                                                                                                                                                                                                                                                                                   TISSUE=Embryo;
                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=22341132; PubMed=12454917; Klein S.L., Strausberg R.L., Wagner
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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(TrEMBLrel. 27,
(TrEMBLrel. 27,
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CU2_MONOOXYGENASE_1;
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DOI=10.1002/dvdy.10174;
r L., Pontius J., Clifton S.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      more than 15,000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              research: The NIH Xenopus
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ID BPT2_BOVIN
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Matches 29
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                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=81044408; PubMed=7428928; DOI=10.1016/0014-5793(80)80997-5; Wachter E., Deppner K., Hochstrasser K., Lempart K., Geiger R.; "A new Kunitz-type inhibitor from bovine serum amino acid sequence determination.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Serum basic protease inhibitor (Serum BPI).
Bos taurus (Bovine).
Eukaryota, Metazoa; Chordara, Craniata, Vertebrata, Euteleostomi,
Mammalia; Eutheria, Cetartiodactyla, Ruminantia, Pecora, Bovidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-JUL-1986 (Rel. 01,
21-JUL-1986 (Rel. 01,
05-JUL-2004 (Rel. 44,
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ProDom; PD000222; Prot_Inh_Kunz-m;
SMART; SM00131; KU; 3.
                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                            PIR; A01206; TIBOR.
HSSP; P00974; 1BPI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR002223; Prot_Inh_Kunz-m.
InterPro; IPR008296; TPFI.
Pfam; PF00014; Kunitz BPT1; 3.
PIRSF; PIRSF001620; TFFI; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P00975;
                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NON TER
                                                                                                                                                                                                                                                                                                                                                                 FEBS Lett. 119:58-62(1980).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IBPS
                                                                                                                                                                                                                                                                       nterPro; IPR002223; Prot Inh Kunz-m. Pfam; PF00014; Kunitz_BPTI; 1.
                                                                                                                                                                                                                                                                                                                     SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Contains 1 BPTI/Kunitz inhibitor domain
                                                                                                                                                                                                                                                                                                                                          FUNCTION: This inhibitor has activity very similar to that basic protease inhibitor from bovine tissues.
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                                                                            PVRCILPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YSEYCAAPSLTGPCRASFSRWYYDTTSGQCATFIYGGCRGNXNNYLSEDDCVKNCVG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQG 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PS00280; BPTI_KUNITZ_1;
PS50279; BPTI_KUNITZ_2;
                                                           PDFCLEPPYTGPCKAAMIRYFYNAKAGFCETFVYGGCRAKSNNFKSAEDCMRTCGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      327 AA; 35951 MW; 2BFE6FA75DAB0497 CRC64;
                                                                                                                                                  60 AA;
                                                                                                       Conservative
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                                                                                                                                                  6647 MW;
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                                                                                                                Score 184; DB 1;
Pred. No. 1.3e-14;
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B9953EBAACF1A4E6 CRC64;
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Pred. No. ;
                                                                                                      Pred. No. 1.30); Mismatches
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similarity.
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                                                                                                                                              MEDLINE-91098258; PubMed=1986787; DOI=10.1016/0167-4838(91)90231-N; Barra D., Fioretti E., Angeletti M., Maras B., Bossa F., Ascoli F.; "Proteinase isoinhibitors from bovine spleen: primary structure of intermediate in the processing of the precursor."; Biochim. Biophys. Acta 1076:143-147(1991).

-i- SUBCELLULAR LOCATION: Secreted.
-i- SIMILARITY: Contains 1 BPTI/Kunitz inhibitor domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=88007630; PubMed=3654647;
Barra D., Simmaco M., Bossa F., Fioretti E., Angeletti M., Ascoli "Primary structure of a protease isoinhibitor from bovine spleen. possible intermediate in the processing of the primary gene production. J. Biol. Chem. 262:13916-13919(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=89076531; PubMed=2462435;
Fioretti E., Angeletti M., Fiorucci L., I
"Aprotinin-like isoinhibitors in bovine o
Biol. Chem. Hoppe-Seyler 369:37-42(1988)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=87283904; PubMed=2441071;
Creighton T.E., Charles I.G.;
"Sequences of the genes and polypeptide precursors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-AUG-1987 (Rel. 05, Created)
01-MAIG-1989 (Rel. 10, Last sequence update)
05-UIL-2004 (Rel. 44, Last annotation update)
Spleen trypsin inhibitor I precursor (SI-I) [Contains: Spleen trypsin inhibitor II (SI-III)].
Bos taunus (Revins); Spleen trypsin inhibitor III (SI-III)].
                                                                                                                                                                                                                                                                                                                         Fioretti B., Iacopino G., Angeletti M., "Primary structure and antiproteolytic inhibitor from bovine spleen.", J. Biol. Chem. 260:11451-11455(1985).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=88295740; PubMed=2456884; Creighton T.E., Charles I.G.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bos taurus (Bovine).
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13-AUG-1987
                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE OF 36-93.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biochem. J. 233:443-450(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Sequences encoding two trypsin inhibitors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Biosynthesis, processing, and evolution of bovine pancreatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u></u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mol. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Spring Harb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inhibitors.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        194:11-22(1987).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  52:511-519(1987).
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activity of a Kunitz-type
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EMBL; AY271744; AAP22956.1; -.
HSSP; Q16019; 1AAP
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CHAIN
DOMAIN
                                             ZFIN; ZDB-GENE-020220-1; appb.
GO; GO:0004867; F:serine-type endopeptidase
InterPro; IPR002223; Prot Inh Kunz-m.
Pfam; PF00014; Kunitz BPTT; 1.
PRINTS; PR00759; BASICFTASE.
Probom; PD000222; Prot Inh Kunz-m; 1.
SMART; SM00131; KU; 1.
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                                                                                                                                                                               Actinopterygii; Neopterygii; Cyprinidae; Danio. CyDrinidae; Danio. CBI_TaxID=7955;
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01-OCT-2003
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01-JUN-2003
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EMBL; X03366; CAA27064.1;
EMBL; X03366; CAA27065.1;
 SEQUENCE
                                                                                                                                          Groth C., Lardelli M.;
Submitted (APR-2003) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                     SEQUENCE FROM N.A.
Groth C., Lardelli
                                                                                                                                                                                                             Brachydanio rerio (Zebrafish)
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                         Amyloid protein b
                                                                                                                                                                                                                                                                                          Q7ZZT3
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PROSITE; PS50279; BPTI_KUNITZ_2; 1.
Direct protein sequencing; Serine p
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SMART; SM00131; KU;
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Pfam; PF00014; Kunitz BPTT; 1.
PRINTS; PR00759; BASICPTASE.
                             PROSITE;
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X06685;
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M20933;
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                           PS00280; BPTI_KUNITZ_1; PS50279; BPTI_KUNITZ_2;
                                                                                                                                                                                                                                                                                                                                        PDFCLEPPYTGPCKAKMIRYFYNAKAGFCETFVYGGCKAKSNNFRSAEDCMRTCGGA
                                                                                                                                                                                                                                                                                                                                                           PVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
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AAA51418.1;
CAA28887.1;
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Spleen trypsin i
Spleen trypsin i
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BPTI/Kunitz inhi
Reactive bond (E
By similarity.
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                                                                                                                                                                                                 ; Craniata; Vertebrata; Euteleostomi; Teleostei; Ostariophysi; Cypriniformes;
                                                                                                                                                                                                                                                                                                                                                                             Score 184; DB 1;
Pred. No. 2.1e-14;
1; Mismatches 16
 0723D4AC6585B349 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rine protease inhibitor; Potential.
                                                                                                                                                                                                                                                                                          PRT;
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                                                                                                                                                                                                                      (Danio rerio)
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                                                                                                                                                                                                                                                                                          82
                                                                                                                                   inhibitor domain
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                                                                                             inhibitor activity;
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Query Match

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RESULT 15
UPTI_PIG
ID UPTI_PIG
AC Q29100;
DT 01-NOV-1997
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Pfam; PR00014; Kunitz BPTI; 1.
PRINTS; PR00759; BASICPTASE.
ProDom; PD000222; Prot Inh Kunz-m; 1.
PROSITE; PS00280; BPTI KUNITZ 1; 1.
PROSITE; PS50279; BPTI KUNITZ 2; 1.
                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collabora between the Swiss Institute of Bioinformatics and the EMBL outstati the European Bioinformatics Institute. There are no restrictions on use by non-profit institutions as long as its content is in no modified and this statement is not removed. Usage by and for commer entities requires a license agreement (See http://www.isb-sib.ch/annou or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Stallings-Mann M.L., Burke M.G., Trout W.E., Roberts R.M.;
"Purification, characterization, and cDNA cloning of a Kunitz-type
proteinase inhibitor secreted by the porcine uterus.";
J. Biol. Chem. 269:24090-24094[1994].
-I- FUNCTION: Inhibitor of plasmin and trypsin. Also has a weak
affinity for chymotrypsin. Colud serves to neutralize the
activities of one or more serine proteinases generated by the
proliferating trophoblast during the formation of the noninvasive
placenta.
-I- TISSUE SPECIFICITY: Expressed only in the uterus and the
                                                                                                                                                                                               SIGNAL
                                                                                                                                                                                                            Direct
                                                                                                                                                                                                                                                                                                                    EMBL; L14282; AAA62425.1; -. PIR; A55115; A55115.
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-!- SIMILARITY: Contains 1 BPTI/Kunitz inhibitor domain.
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Mammalia; Eutheria;
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01-NOV-1997 (Rel. 35, Last sequence update)
05-UUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sus scrofa (Pig)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Uterine plasmin/trypsin inhibitor precursor (UPTI).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DEVELOPMENTAL STAGE: Maximally expressed during pregnancy until day 30 after which levels decrease significantly.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                endometrium.
                                                                                                                                                                                                                                                                                                          P00974; 1K09
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29; Conservative
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                            sequencing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
                                          SEQUENCE OF 30-56, AND CHARACTERIZATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chordata; Craniata; Vertebrata; Euteleostomi; Cetartiodactyla; Suina; Suidae; Sus.
                            13109 MW;
  52.9%;
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                        Uterine plasmin/trypsin inhibitor.
BFTI/Kunitz inhibitor.
Reactive bond (By similarity).
By similarity.
By similarity.
G -> A (in Ref. 1; AA sequence).
S -> R (in Ref. 1; AA sequence).
V -> I (in Ref. 1; AA sequence).
V -> V (in Ref. 1; AA sequence).
Y -> V (in Ref. 1; AA sequence).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and Uterus;
                                                                                                                                                                                            Serine protease inhibitor; Signal. Potential.
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7; Mismatches 18; Indels
  Score 174;
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Length 122;
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MBL outstation -
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<u>,,</u>

Search completed: August 25, 2005, 04:55:26
Job time : 180 secs

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Result
No.
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Maximum Match
Listing first
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Perfect score:
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  Score
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seg length: 2000000000
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Query
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... /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
... /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
... /cgn2_6/ptodata/1/pubpaa/USO6_PUBCOMB.pep:*
... /cgn2_6/ptodata/1/pubpaa/USO6_PUBCOMB.pep:*
... /cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
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... /cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
... /cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
... /cgn2_6/ptodata/1/pubpaa/USO9A_PUBCOMB.pep:*
... /cgn2_6/ptodata/1/pubpaa/USO9C_PUBCOMB.pep:*
... /cgn2_6/ptodata/1/pubpaa/USOC_PUBCOMB.pep:*
... /cgn2_6/ptodata/1/pubpaa/USOOC_PUBCOMB.pep:*
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Length
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140.197 Million cell updates/sec
Description
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ALIGNMENTS

US-09-896-322-5
US-09-234-874A-7
US-09-234-874A-7
US-09-234-874A-7
US-09-234-874A-7
US-09-113-244-2
US-09-781-988-44
US-09-781-988-44
US-09-781-988-6-05-44
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US-09-896-095-144
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US-10-778-986-1

Sequence 86, Appli
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Sequence 108, App
Sequence 108, App
Sequence 61, Appli
Sequence 2, Appli
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Sequence 44, Appli
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ntive 0	Query Match 100.0%; Score 329; DB 16; Length 58; Best Local Similarity 100.0%; Pred. No. 5.6e-32;	; ORGANISM: Homo sapiens US-10-721-961-4	; SEQ ID NO 4 ; LENGTH: 58 ; TYPE: PRT	; NUMBER OF SEQ ID NOS: 34 ; SOFTWARE: PatentIn version 3.2	PRIOR FILING DATE: 2002-05-31	; PRIOR APPLICATION NUMBER: US 60/303,180	; PRIOR FILING DATE: 2001-05-31	; CURRENT FILING DATE: 2003-11-25 ; PRIOR APPLICATION NUMBER: Danish Application No. PA 2001 00859	CURRENT APPLICATION NUMBER: US/10/721,961	FILE REFERENCE: 6297.204-US	; TITLE OF INVENTION: Kunitz-Type Sequences and Polypeptides		; APPLICANT: Susanne, Bang	; APPLICANT: Jorgensen, Marianne U	; APPLICANT: Novo Nordisk Pharmaceuticals, Inc.	; GENERAL INFORMATION:	; Publication No. US20040152633A1	; Sequence 4, Application US/10721961	US-10-721-961-4	RESULT 1	

100.0 99.11 95.4 88.4 88.4 80.2 80.2 80.2

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YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58

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APPLICANT: Petersén, Lars C
ITITLE OF INVENTION: Kunitz-Type Sequences and Polypeptides
FILE REFERENCE: 6297.204-US
CURRENT APPLICATION NUMBER: US/10/721,961
CURRENT FILING DATE: 2003-11-25
PRIOR APPLICATION NUMBER: Danish Application No. PA 2001 00859
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: US 60/303,180
PRIOR APPLICATION NUMBER: US 60/303,180
PRIOR APPLICATION NUMBER: PCT/DK02/00372
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US-10-721-961-5
                                                                                                                                                                                                      ; SEQ ID NO 7
; LENOTH: 58
; TYPE: PRT
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-721-961-7
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                                                                         Matches
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Publication No. US20040152633A1
                                                                                                     Best Local
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Petersen, Lars C
TITLE OF INVENTION: Kunitz-Type Sequences and Polypeptides
FILE REFERENCE: 6297.204-US
CURRENT APPLICATION NUMBER: US/10/721,961
CURRENT FILING DATE: 2003-11-25
PRIOR APPLICATION NUMBER: Danish Application No. PA 2001 00859
PRIOR PILING DATE: 2001-05-31
PRIOR PILING DATE: 2001-07-05
PRIOR APPLICATION NUMBER: US 60/303,180
PRIOR PILING DATE: 2001-07-05
PRIOR APPLICATION NUMBER: PCT/DK02/00372
                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 2002-05-31 NUMBER OF SEQ ID NOS: 34
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APPLICANT: Jorgensen, Marianne U
APPLICANT: Susanne, Bang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Novo Nordisk Pharmaceuticals, APPLICANT: Jorgensen, Marianne U
                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.2
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APPLICANT: Olesen, Ole H
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TYPE: PRT
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                                                                     Similarity 55; Conserv
1 YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
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Olesen, Ole H
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                                                                                              95.4%; Score 314; DB 16; 94.8%; Pred. No. 3.6e-30;
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                                                                     Mismatches
                                                                                                                                 Length 58;
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APPLICANT: Petersen, Lars C
TITLE OF INVENTION: Kunitz-Type Sequences and P(
FILE REPERENCE: 6297.204-US
CURRENT APPLICATION NUMBER: US/10/721,961
CURRENT FILING DATE: 2003-11-25
PRIOR APPLICATION NUMBER: Danish Application No.
PRIOR TILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: US 60/303,180
PRIOR APPLICATION NUMBER: US 60/303,180
PRIOR APPLICATION NUMBER: PCT/DK02/00372
PRIOR FILING DATE: 2001-07-05
PRIOR FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 34
                                                                 ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-721-961-6
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US-10-721-961-34
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Best Local Similarity
Matches 55; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn version 3.2 SEQ ID NO 34
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Query Match
Best Local Similarity
                                                                                                                                                       SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/721,961
CURRENT FILING DATE: 2003-11-25
PRIOR APPLICATION NUMBER: Danish Application |
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: US 60/303,180
PRIOR PRIOR PRIOR DATE: 2001-07-05
PRIOR PRIOR APPLICATION NUMBER: US 7-05-31
PRIOR PRIOR DATE: 2002-05-31
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Petersen, Lars C
TITLE OF INVENTION: Kunitz-Type Sequences
FILE REFERENCE: 6297.204-US
                                                                                                                                                                         SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                              NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Novo Nordisk Pharmaceuticals, Inc
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ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                              ENGTH: 58
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: Susanne, Bang
: Olesen, Ole H
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Olesen, Ole H
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Vo. US20040152633A1
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  88.4%;
89.7%;
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Pred. No. 6.6e-30;
0; Mismatches 3;
  Score 291; DB 16;
Pred. No. 2.1e-27;
                                                                                                                                                                                                                                                                                                                                                                                                                    and Polypeptides
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                                                                                                                                                                                                                                                                                                                                 No.
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                  Length 58;
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FILE REFERENCE: 6297.204-US
CURRENT APPLICATION NUMBER: US/10/721,961
CURRENT FILING DATE: 2003-11-25
PRIOR APPLICATION NUMBER: DATISH Application No.
PRIOR PILING DATE: 2001-05-31 *
PRIOR APPLICATION NUMBER: US 60/303,180
PRIOR FILING DATE: 2001-07-05,*
PRIOR APPLICATION NUMBER: PCT/DK02/00372
PRIOR APPLICATION NUMBER: PCT/DK02/00372
PRIOR APPLICATION NUMBER: PCT/DK02/00372
PRIOR PILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn version 3.2
SEQ ID NO 3
SEQ ID NO 3
LENGTH: 111
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                   SEQ ID NO 1
LENGTH: 58
TYPE: PRT
ORGANISM: Artificial
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Publication No. US20040152633A1
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                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/721,961
CURRENT FILING DATE: 2003-11-25
PRIOR APPLICATION NUMBER: Danish Application
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: US 60/303,180
PRIOR APPLICATION NUMBER: US 60/303,180
PRIOR FILING DATE: 2001-07-05
                                                                                                 PRIOR APPLICATION NUMBER: PCT/DK02/00372
PRIOR FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 34
SOFTMARE: PatentIn version 3.2
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TITLE OF INVENTION: Kunitz-Type Sequences
FILE REFERENCE: 6297.204-US
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Novo Nordisk Pharmaceuticals, Inc
APPLICANT: Jorgensen, Marianne U
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APPLICANT: Jorgensen, Marianne U
APPLICANT: Susanne, Bang
APPLICANT: Olesen, Ole H
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TITLE OF INVENTION: Kunitz-Type Sequences and Polypeptides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial
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FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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Olesen, Ole H
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No. US20040152633A1
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Pred. No. 3.
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.8e-27;
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Matches
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SEQ ID NO 32
LENGTH: 111
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                                                                                                                                                                                                                                                                                                            Sequence 92, Application US/102 Publication No. US20040038877A1
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Publication No. US20040152633A1
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TITLE OF INVENTION: Kunitz-Type Sequences and Polypeptides
FILE REFERENCE: 6297.204-US
CURRENT APPLICATION NUMBER: US/10/721,961
CURRENT FILING DATE: 2003-11-25
PRIOR APPLICATION NUMBER: Danish Application No. PA 2001 00859
PRIOR FILING DATE: 2001-05-31
PRIOR PILING DATE: 2001-07-05
PRIOR FILING DATE: 2001-07-05
PRIOR FILING DATE: 2001-07-05
PRIOR PILING DATE: 2001-07-05
PRIOR APPLICATION NUMBER: PCT/DK02/00372
PRIOR APPLICATION STEEL 2001-05-31
NUMBER: OF SEQ ID NOS: 34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Amino acid sequence of the 212L-HKI18 fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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48; Conservative
                                                                                                                                                                                                                                       Alsobrook, John,
Anderson, David W.,
Boldog, Ferenc,
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                                                                                                                                          Ellerman, Karen,
Gerlach, Valerie,
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Olesen, Ole
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Reiger, Daniel,
                          Li, Li,
Miller, Charles,
                                                                                                                                   Gorman,
                                                                                                                                                                                                                       Burgess, Catherine,
              Patturajan, Meera,
                                                              Kekuda, Ramesh,
Leach, Martin,
                                                                                                                                                                                     Edinger, Shlomit,
                                                                                                                                                                                                       Catterton,
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                                                                                                  Weizhen,
                                                                                                                 Xiaojia,
                                                                                                                                   Linda,
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Pred. No. 6.8e-24;
0; Mismatches 10
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1.7e-24;
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; LENGTH: 1280
; TYPE: PRT
; ORGANISM: Homo &
US-10-262-839-92
                                    us-10-103-377C-6
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US-10-103-377C-6
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                                                                                                                                                                                                                                                                                                                 Sequence 6, Application US/10103377C Publication No. US20030073098A1 GENERAL INFORMATION:
 Query Match
                                                                                                              SOFTWARE: FASTSEQ
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: CO
SEQ ID NO 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                       LENGTH: 1311
TYPE: PRT
ORGANISM: Homo
                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/103,377C CURRENT FILING DATE: 2002-08-26 PRIOR APPLICATION NUMBER: 60/278,347 PRIOR FILING DATE: 2001-03-23 NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                           APPLICANT: Bandaru, Rajasekhar
TITLE OF INVENTION: 65577, A Human Matrix
TITLE OF INVENTION: Metalloproteinase and Uses
FILE REFERENCE: MPI01-049P1RNM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/326,483
PRIOR FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: 60/327,917
PRIOR FILING DATE: 2001-10-09
PRIOR APPLICATION NUMBER: 60/328,029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2001-10-09
PRIOR APPLICATION NUMBER: 60/328,056
PRIOR FILING DATE: 2001-10-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Zerhusen, Brian APPLICANT: Zhong, Mei TITLE OF INVENTION: THERAPIFILE REFERENCE: 21402-462A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/381,101
PRIOR FILING DATE: 2002-05-16
PRIOR APPLICATION NUMBER: 60/371,972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/262,839
CURRENT FILING DATE: 2002-10-01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 2001-10-09
APPLICATION NUMBER: 60/328,849
FILING DATE: 2001-10-12
APPLICATION NUMBER: 60/374,738
FILING DATE: 2002-04-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/327,342
FILING DATE: 2001-10-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/328,044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CuraSeqList version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                    YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 809
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Taupier, koj.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Spytek, Kimberly,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Smithson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rothenberg, Mark,
Shimkets, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zerhusen, Brian,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Voss, Edward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                       sapiens
                                                                                                                                 for Windows Version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 THERAPEUTIC POLYPEPTIDES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Raymond,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Glennda,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Richard,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           80.2%;
   80.2%;
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   Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 264; DB 15;
Pred. No. 6.7e-23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
   264;
                                                                                                                                 4.0
 В
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   14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10;
                                                                                                                                                                                                                                                                Therefor
Length 1311;
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                                                                         Query Match
Best Local S
Matches 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 91, Application US/103 Publication No. US20040121349A1
                                                                                                                                                                                                                                   SEQ ID NO 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 82.0 Matches 48; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 09/950,370
PRIOR FILING DATE: 2001-09-10
PRIOR APPLICATION NUMBER: US 60/231,084
PRIOR FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: US 10/294,039
PRIOR FILING DATE: 2002-11-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: MPI03-0190MNIM
CURRENT APPLICATION NUMBER: US/10/391,364
CURRENT FILING DATE: 2003-03-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bandaru, Rajasekhar
TITLE OF INVENTION: NOVEL 27877, 18080,
TITLE OF INVENTION: 16588, 14223, 1600:
TITLE OF INVENTION: USES THEREFOR
                                                                                                                                                                                                TYPE: PR
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                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/327,820 PRIOR FILING DATE: 2001-10-09
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                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 09/717,926
FILING DATE: 2000-11-21
APPLICATION NUMBER: US 60/214,707
FILING DATE: 2000-06-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 10/266,035 FILING DATE: 2002-10-07 APPLICATION NUMBER: US 60/328,198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 60/338,587 FILING DATE: 2001-11-13
                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 10/268,036 FILING DATE: 2002-10-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 2001-10-09
732
                                                                                                                                                                                                              1311
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                                                                                            Similarity
                                                                                                                                                                                                                                                       FastSEQ for Windows Version 4.0
YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 789
                                    YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 789
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kapeller-Libermann,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cook, William James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Carroll,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/10391364
                                                                           Conservative
                                                                                            80.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           82.8%;
                                                                         Score 264; DB 16;
Pred. No. 6.8e-23;
0; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ,
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D; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rosana
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16002, 50566, 65552 AND 65577 MOLECULES
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                                                                                                                                                                                                                                                                                            See
                                                                                                                                                                                                                                                                                          File Wrapper
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                                                                                                              Length 1311;
                                                                           Indels
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                                                                         Gaps
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; Sequence 86, Application US/10087887
; Publication No. US20030.98957A1
; GENERAL INFORMATION:
; APPLICANT: Kekuda, Ramesh
; APPLICANT: Conley, Pamela B.
; APPLICANT: Yang, Ruey-Bing
; APPLICANT: Hart, Matthew

RESULT 12 US-10-087-887-86

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; TYPE: PRT
; ORGANISM: Mus musculus
US-10-087-887-86
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 5, Application US/08896322
Publication No. US20020103334A1
GENERAL INFORMATION:
APPLICANT: Werner Schroder, Sorv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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TITLE OF INVENTION: PROTEINS AND NUCLEIC ACIDS ENCODING SAME
FILE REFERENCE: 21402-285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/087,887
CURRENT FILING DATE: 2002-03-01
CURRENT PRIOR APPLICATION NUMBER: 60/273,049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 104
SOFTWARE: CuraSeqList version 0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                   FILING DATE: 17-JUL-1997
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION UNDATA:
APPLICATION UNDATA: 196 29 982.9
FILING DATE: 25-JUL-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
COMPUTER: HP VECTRA
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Werner Schroder, Soren Bjorn, Kjeld No. US20020103334A1ris,Viggo Diness
TITLE OF INVENTION: APROTININ VARIANTS HAVING IMPROVED PRO
NUMBER OF SEQUENCES: 5
                                                                                                                                                                              SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                             STREET: 600
CITY: Tarrytown
                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
NAME: Kurt G. Briscoe
REGISTRATION NUMBER: 33
REFERENCE/DOCKET NUMBER:
                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                    OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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FILING DATE: 2001-04-03
APPLICATION NUMBER: 60/282,864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/282,537 FILING DATE: 2001-04-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/279,883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 2001-03-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/282,867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 2001-04-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 2001-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       746 YPVRCLLPSAQGSCGDWAARWYFVASVGRCNRFWYGGCHGNANNFASEQECMNTCRG 802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQG 57
                                                                                                                                                                                                                                                                                            10591-5144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zerhusen, Bryan
Komuves, Laszlo
                                                                                                                                                                                                                                                                                                                                                                   660 White Plains Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shimkets, Richard A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tomlinson,
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                                                                                                                                                                                                                                                                                                              U.S.A.
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en, Bryan D.
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                                                                                                                                                                                                                    DOS
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75.4%;
                                                                                                                                                                                                                                                                                                                                                                                KRAMER SCHAEFER & BRISCOE
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                                                                                                                                                               US/08/896,322
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                                                                                                                                                                                                                                                      3.50 inch,
 Bayer 9911-KGB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 246; DB 14;
Pred. No. 9.8e-21;
4: Mismatches 10;
                                                                                       (Germany)
                                                                                                                                                                                                                                                     2.0 MB storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1280;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
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JEQUENCE DESCRIPTION: SEQ ID NO: US-10-252-967-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-252-967-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-896-322-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/10252967 Publication No. US20030096752A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                             TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: Peptide FRAGMENT TYPE: linear ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 57 Amino acids
TYPE: Amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (914) 332-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (914) 332-170
TELEFAX: (914) 332-1844
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/896,322
FILING DATE: 17-UUL-1997
ATTORNEY/AGENT INFORMATION:
NAME: Kurt G. Briscoe
REGISTRATION NUMBER: 33,141
                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: Bayer 9911-KGB TELECOMMUNICATION INFORMATION: TELEPHONE: (914) 332-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/252,967
                                                                    ORIGINAL SOURCE:
                                                                                   MOLECULE TYPE: FRAGMENT TYPE:
                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Werner Schroder, Soren Bjorn, Kjeld No. US20030096752Alris, Viggo Dir
TITLE OF INVENTION: APROTININ VARIANTS HAVING IMPROVED PRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 CLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                              TYPE: Amino acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: HP VECTRA
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: New York COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLEPPSTGPCRAAIIRYFYDATAGLCETFVYGGCRANRNNFKSAEDCMETCGGA 57
                                                                                                                             TOPOLOGY:
                                                                                                                                                                                       LENGTH: 57 Amino acids
                                                                                                                                                                                                                                                                     TELEFAX: (914) 332-1844
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 23-Sep-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: WordPerfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Diskette, COMPUTER: HP VECTRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Tarrytown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: SPRUNG KRAMER SCHAEFER & BRISCOE STREET: 660 White Plains Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10591-5144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aprotinin variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Single
                                                                                                      Peptide
                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     58.7%; Score 193; DB 8; 57.4%; Pred. No. 1.2e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>ب</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.50 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.0 MB storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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TELEX: 904136
; INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 58 amino acids
TYPE: amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-09-234-874A-7
Search completed: August 25, 2005, 04:59:08 Job time : 164 secs
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US-09-234-874A-7
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Best Local Similarity
Matches 33; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7, Application US/09234874A Publication No. US20010020003A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: D.C.

COUNTRY: USA
ZIP: 20007-5109
ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/234,874A
PILING DATE: 11-Jun-2001
PRIOR APPLICATION UNMBER: US/09/234,874A
FILING DATE: 08-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Bent, Stephen
REGISTRATION NUMBER: 09/768
REFERENCE/DOCKET NUMBER: 056324/0106
TELEPAN: (202)672-5309
TELEPAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lesikar, David D.
McFadden, Kathleen
Garrick, Brett L.
TITLE OF INVENTION: PROTEASE INHIBITOR PEPTIDES
NUMBER OF SEQUENCES: 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: White, Tyler R. Damm, Deborah
                                                                                                                        2 PVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5 CLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
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                                                                                         PDFCLEPPYTGPCKARIIRYFYNAKAGLCQTFVYGGCRAKRNNFKSAEDCMRTCGGA 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Washington
                                                                                                                                                                                      Conservative
                                                                                                                                                                               58.4%; Score 192; DB 9; Length 58; 57.9%; Pred. No. 1.7e-15; ative 7; Mismatches 17; Indels
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